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(57) Abstract

A method for protecting a peptide from peptidase activity in vivo, the peptide being composed of between 2 and 50 amino acids and having a C-terminus and an N-terminus and a C-terminus amino acid and an N-terminus amino acid is described. In the first step of the method, the peptide is modified by attaching a reactive group to the C-terminus amino acid, to the N-terminus amino acid, or to an amino acid located between the N-terminus and the C-terminus, such that the modified peptide is capable of forming a covalent bond in vivo with a reactive functionality on a blood component. In the next step, a covalent bond is formed between the reactive group and a reactive functionality on a blood component to form a peptide-blood component conjugate, thereby protecting said peptide from peptidase activity. The final step of the method involves the analyzing of the stability of the peptide-blood component conjugate to assess the protection of the peptide from peptidase activity.

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PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD COMPONENTS

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FIELD OF THE INVENTION

This invention relates to modified therapeutic peptides. In particular, this invention relates to protection of endogenous therapeutic peptides from peptidase activity through a modification that enables the peptide to selectively conjugate to blood components, thus protecting the peptide from peptidase activity and increasing the duration of action of the therapeutic peptide for the treatment of various disorders.

BACKGROUND OF THE INVENTION

Many endogenous peptides have been described as key components of biological processes. Some of these peptides have been identified as key therapeutic agents for the management of various disorders. In general, endogenous peptides are more desirable as therapeutic agents than synthetic peptides with non-native sequences, because they do not produce an immune response due to their endogenous character. In addition, endogenous peptides are highly specific for their target receptors and are easy to synthesize and manufacture. However, a major difficulty with the delivery of such therapeutic peptides is their short plasma half-life, mainly due to rapid serum clearance and proteolytic degradation via the action of peptidases.

Peptidases break a peptide bond in peptides by inserting a water molecule across the bond. Generally, most peptides are broken down by peptidases in the body in a manner of a few minutes or less. In addition, some peptidases are specific for certain types of peptides, making their degradation even more rapid. Thus, if a peptide is used as a therapeutic agent, its activity is generally reduced as the peptide quickly degrades in the body due to the action of peptidases.

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One way to overcome this disadvantage is to administer large dosages of the therapeutic peptide of interest to the patient so that even if some of the peptide is degraded, enough remains to be therapeutically effective. However, this method is quite uncomfortable for the patient.

Since most therapeutic peptides cannot be administered orally, the therapeutic peptide would have to be either constantly infused, frequently administered by intravenous injections, or administered frequently by the inconvenient route of subcutaneous injections. The need for frequent administration also results in many potential peptide therapeutics having an unacceptably high projected cost per treatment course. The presence of large amounts of degraded peptide may also generate undesired side effects.

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Discomfort in administration and high costs are two reasons why most therapeutic peptides with attractive bioactivity profiles are not developed as drug candidates. Instead, these therapeutic peptides are used as templates for the development of peptidomimetic compounds to substitute for the therapeutic peptide. Biotechnology and large pharmaceutical firms frequently undertake lengthy and expensive optimization programs to attempt to develop non-peptide, organic compounds which mimic the activity seen with therapeutic peptides without incurring an unacceptable side effect profile. For example, cyclic peptides, peptidomimetics and small molecules coming from expensive SAR (Structure Activity Relationship) and molecular modeling studies have led to the development of an incredible amount of peptide mimics. However, these peptide mimics in no way reflect the exact original biological nature of the therapeutic peptide, and thus are inferior to the endogenous therapeutic peptide as therapeutic agents.

An alternative to creating peptide mimics is to block the action of peptidases to prevent degradation of the therapeutic peptide or to modify the therapeutic peptides in such a way that their degradation is slowed down while still maintaining biological activity. Such methods include conjugation with polymeric materials such as dextrans, polyvinyl

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pyrrolidones, glycopeptides, polyethylene glycol and polyamino acids, conjugation with adroitin sulfates, as well as conjugation with polysaccharides, low molecular weight compounds such as aminolethicin, fatty acids, vitamin B₁₂, and glycosides. These conjugates, however, are still often susceptible to protease activity. In addition, the therapeutic activity of these peptides is often reduced by the addition of the polymeric material. Finally, there is the risk of the conjugates generating an immune response when the material is injected *in vivo*. Several methods include *ex vivo* conjugation with carrier proteins, resulting in the production of randomized conjugates. Since conjugates are difficult to manufacture, and their interest is limited by commercial availability of the carriers, as well as by their poor pharmaco economics.

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There is thus a need for novel methods to modify therapeutic peptides to protect them from peptidase activity and to provide longer duration of action *in vivo*, while maintaining low toxicity yet retaining the therapeutic advantages of the modified peptides.

SUMMARY OF THE INVENTION

This invention is directed to overcoming the problem of peptide degradation in the body by modifying the therapeutic peptide of interest and attaching it to protein carriers, such that the action of peptidases is prevented, or slowed down. More specifically, this invention relates to novel chemically reactive derivatives of therapeutic peptides that can react with available functionalities on blood proteins to form covalent linkages, specifically a therapeutic peptide-maleimide derivative. The invention also relates to novel chemically reactive derivatives or analogs of such therapeutic peptides. The invention additionally pertains to the therapeutic uses of such compounds.

The present invention is directed to modifying and attaching therapeutic peptides to protein carriers, preferentially albumin, through *in vivo* or *ex vivo* technology to prevent or reduce the action of peptidases

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by virtue of a synthetic modification on the first residue to be cleaved. Therapeutic peptides are usually active at the N-terminus portion, at the C-terminus portion, or in an interior portion of the peptide chain. Using the technology of this invention, a site other than the active portion of a therapeutic peptide is modified with certain reactive groups. These reactive groups are capable of forming covalent bonds with functionalities present on blood components. The reactive group is placed at a site such that when the therapeutic peptide is bonded to the blood component, the peptide retains a substantial proportion of the parent compound's activity.

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The modification of the therapeutic peptide through the chemical modification used in the invention is done in such a way that all or most of the peptide specificity is conserved despite attachment to a blood component. This therapeutic peptide-blood component complex is now capable of traveling to various body regions without and being degraded by peptidases, with the peptide still retaining its therapeutic activity. The invention is applicable to all known therapeutic peptides and is easily tested under physiological conditions by the direct comparison of the pharmacokinetic parameters for the free and the modified therapeutic peptide.

The present invention is directed to a modified therapeutic peptide capable of forming a peptidase stablilized therapeutic peptide composed of between 3 and 50 amino acids. The peptide has a carboxy terminal amino acid, an amino terminal amino acid, a therapeutically active region of amino acids and a less therapeutically active region of amino acids. The peptide comprises a reactive group which reacts with amino groups, hydroxyl groups, or thiol groups on blood components to form a stable covalent bond and thereby forms the peptidase stabilized therapeutic peptide. In the peptide of the invention the reactive group is selected from the group consisting of succinimidyl and maleimido groups and the reactive group is attached to an amino acid positioned in the less therapeutically active region of amino acids.

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In one embodiment, the therapeutically active region of the peptide includes the carboxy terminal amino acid and the reactive group is attached to said amino terminal amino acid.

In another embodiment, the therapeutically active region of the peptide includes the amino terminal amino acid and the reactive group is attached to the carboxy terminal amino acid.

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In another embodiment, the therapeutically active region of the peptide includes the carboxy terminal amino acid and the reactive group is attached to an amino acid positioned between the amino terminal amino acid and the carboxy terminal amino acid.

In yet another embodiment, the therapeutically active region includes the amino terminal amino acid and the reactive group is attached to an amino acid positioned between the amino terminal amino acid and the carboxy terminal amino acid.

The present invention is also directed to a method of synthesizing the modified therapeutic peptide. The method comprises the following steps. In the first step, if the therapeutic peptide does not contain a cysteine, then the peptide is synthesized from the carboxy terminal amino acid and the reactive group is added to the carboxy terminal amino acid. Alternatively, a terminal lysine is added to the carboxy terminal amino acid and the reactive group is added to the terminal lysine. In the second step, if the therapeutic peptide contains only one cysteine, then the cysteine is reacted with a protective group prior to addition of the reactive group to an amino acid in the less therapeutically active region of the peptide. In the third step, if the therapeutic peptide containe two cysteines as a disulfide bridge, then the two cysteines are oxidized and the reactive group is added to the amino terminal amino acid, or to the carboxy terminal amino acid, or to an amino acid positioned between the carboxy terminal amino acid and the amino terminal amino acid of the therapeutic peptide. In the fourth step, if the therapeutic peptide contains more than two cysteines as disulfide bridges, the cysteines are sequentially oxidized in the disulfide bridges

and the peptide is purified prior to the addition of the reactive groups to the carboxy terminal amino acid.

The present invention is also directed to a method for protecting a therapeutic peptide from peptidase activity in vivo, the peptide being composed of between 3 and 50 amino acids and having a carboxy terminus and an amino terminus and a carboxy terminal amino acid amino acid and an amino terminal amino acid. The method comprises the following steps:

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- (a) modifying the peptide by attaching a reactive group to the carboxy terminal amino acid, to the amino terminal amino acid, or to an amino acid located between the amino terminal amino acid and the carboxy terminal amino acid, such that the modified peptide is capable of forming a covalent bond in vivo with a reactive functionality on a blood component;
- (b) forming a covalent bond between the reactive group and a reactive functionality on a blood component to form a peptide-blood component conjugate, thereby protecting the peptide from peptidase activity; and
- (c) analyzing the stability of the peptide-blood component conjugate to assess the protection of the peptide from peptidase activity. These steps may be performed either *in vivo* or *ex vivo*.

The present invention is also directed to a method for protecting a therapeutic peptide from peptidase activity in vivo, the peptide being composed of between 3 and 50 amino acids and having a therapeutically active region of amino acids and a less therapeutically active region of amino acids. The method comprises the following steps:

- (a) determining the therapeutically active region of amino acids:
- (b) modifying the peptide at an amino acid included in the less
 30 therapeutically active region of amino acids by attaching a reactive
 group to the amino acid to form a modified peptide, such that the

modified peptide has therapeutic activity and is capable of forming a covalent bond in vivo with a reactive functionality on a blood component;

- (c) forming a covalent bond between the reactive entity and a
 reactive functionality on a blood component to form a peptide-blood
 component conjugate, thereby protecting the peptide from peptidase
 activity; and
 - (d) analyzing the stability of the peptide-blood component conjugate to assess the protection of the peptide from peptidase activity. These steps may be performed either *in vivo* or *ex vivo*.

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The peptides useful in the compositions and methods of the present invention include, but are not limited to, the peptides presented in SEQ ID NO:1 to SEQ ID NO:1617.

DETAILED DESCRIPTION OF THE INVENTION

DEFINITIONS

To ensure a complete understanding of the invention, the following definitions are provided:

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Reactive Groups: Reactive groups are entities capable of forming a covalent bond. Such reactive groups are coupled or bonded to a therapeutic peptide of interest. Reactive groups will generally be stable in an aqueous environment and will usually be carboxy, phosphoryl, or convenient acyl group, either as an ester or a mixed anhydride, or an imidate, thereby capable of forming a covalent bond with functionalities such as an amino group, a hydroxy or a thiol at the target site on mobile blood components. For the most part, the esters will involve phenolic compounds, or be thiol esters, alkyl esters, phosphate esters, or the like. Reactive groups include succimidyl and maleimido groups.

Functionalities: Functionalities are groups on blood components, including mobile and fixed proteins, to which reactive groups on modified therapeutic peptides react to form covalent bonds. Functionalities usually include hydroxyl groups for bonding to ester reactive groups, thiol groups for bonding to maleimides, imidates and thioester groups; amino groups for bonding to activated carboxyl, phosphoryl or any other acyl groups on reactive groups.

Blood Components: Blood components may be either fixed or mobile. Fixed blood components are non-mobile blood components and include tissues, membrane receptors, interstitial proteins, fibrin proteins, collagens, platelets, endothelial cells, epithelial cells and their associated membrane and membraneous receptors, somatic body cells, skeletal and smooth muscle cells, neuronal components, osteocytes and osteoclasts and all body tissues especially those associated with the circulatory and lymphatic systems. Mobile blood components are blood components that do not have a fixed situs for any extended period of time, generally not exceeding 5, more usually one minute. These blood components are not membrane-associated and are present in the blood for extended periods of time and are present in a minimum concentration of at least 0.1 μ g/ml. Mobile blood components include serum albumin, transferrin, ferritin and immunoglobulins such as IgM and IgG. The half-life of mobile blood components is at least about 12 hours.

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Protective Groups: Protective groups are chemical moieties utilized to protect peptide derivatives from reacting with themselves. Various protective groups are disclosed herein and in U.S. 5,493,007 which is hereby incorporated by reference. Such protective groups include acetyl, fluorenylmethyloxycarbonyl (Fmoc), t-butyloxycarbonyl (Boc), benzyloxycarbonyl (Cbz), and the like. The specific protected amino acids are depicted in Table 1.

Linking Groups: Linking groups are chemical moieties that link or connect reactive groups to therapeutic peptides. Linking groups may comprise one or more alkyl groups, alkoxy group, alkenyl group, alkynyl group or amino group substituted by alkyl groups, cycloaikyl group, polycyclic group, aryl groups, polyaryl groups, substituted aryl groups, heterocyclic groups, and substituted heterocyclic groups. Linking groups may also comprise poly ethoxy aminoacids such as AEA ((2-amino) ethoxy acetic acid) or a preferred linking group AEEA ([2-(2-amino)ethoxy)]ethoxy acetic acid). A preferred linking group is aminoethoxyethoxyacetic acid (AEEA).

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Sensitive Functional Groups — A sensitive functional group is a group of atoms that represents a potential reaction site on a therapeutic peptide. If present, a sensitive functional group may be chosen as the attachment point for the linker-reactive group modification. Sensitive functional groups include but are not limited to carboxyl, amino, thiol, and hydroxyl groups.

Modified Therapeutic Peptides — A modified therapeutic peptide peptide is a therapeutic peptide that has been modified by attaching a reactive group, and is capable of forming a peptidase stabalized peptide through conjugation to blood components. The reactive group may be attached to the therapeutic peptide either via a linking group, or optionally without using a linking group. It is also contemplated that one or more additional amino acids may be added to the therapeutic peptide to facilitage the attachment of the reactive group. Modified peptides may be administered *in vivo* such that conjugation with blood components occurs *in vivo*, or they may be first conjugated to blood components *in vitro* and the resulting peptidase stabalized peptide (as defined below) administered *in vivo*. The terms "modified therapeutic peptide" and "modified peptide" may be used interchangeably in this application.

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Peptidase Stabalized Therapeutic Peptides - A peptidase stabalized therapeutic peptide is a modified peptide that has been conjugated to a blood component via a covalent bond formed between the reactive group of the modified peptide and the functionalities of the blood component, with or without a linking group. Peptidase stabalized peptides are more stable in the presence of peptidases in vivo than a non-stabalized peptide. A peptidase stabalized therapeutic peptide generally has an increased half life of at least 10-50% as compared to a non-stabalized peptide of identical sequence. Peptidase stability is determined by comparing the half life of the unmodified therapeutic peptide in serum or blood to the half life of a modified counterpart therapeutic peptide in serum or blood. Half life is determined by sampling the serum or blood after administration of the modified and non-modified peptides and determining the activity of the peptide. In addition to determining the activity, the length of the therapeutic peptide may also be measured.

Therapeutic Peptides – As used in this invention, therapeutic peptides are amino acid chains of between 2-50 amino acids with therapeutic activity, as defined below. Each therapeutic peptide has an amino terminus (also referred to as N-terminus or amino terminal amino acid), a carboxyl terminus (also referred to as C-terminus terminal carboxyl terminal amino acid) and internal amino acids located between the amino terminus and the carboxyl terminus. The amino terminus is defined by the only amino acid in the therapeutic peptide chain with a free α -amino group. The carboxyl terminus is defined by the only amino acid in the therapeutic peptide chain with a free α -carboxyl group.

Therapeutic peptides used in the present invention contain a therapeutically active region generally located at the amino terminus, at the carboxyl terminus, or at an internal amino acid. The therapeutically active region may be identified using blind or structure activity

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relationship (SAR) driven substitution, as defined in more detail in this application. SAR is an analysis which defines the relationship between the structure of a molecule and its pharmacological activity for a series of compounds. Alternatively, where the therapeutically active region has previously been defined and is available in the literature, it may be obtained by referring to references such as scientific journals. Knowledge of the location of the therapeutically active region of the peptide is important for modifying the therapeutic peptide, as defined in more detail below.

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Therapeutic peptides used in this invention also contain a less therapeutically active region generally located at the amino terminus, at or near the carboxyl terminus, or at or near an internal amino acid. The less therapeutically active region is a region of amino acids that does not coincide with the therapeutically active region of the therapeutic peptide. The less therapeutically active reion is generally located away from the therapeutically active region, such that modification at the less therapeutically active region does not substantially affect the therapeutic activity of the therapeutic peptide. For example, if the therapeutically active region is located at the amino terminus, the therapeutic peptide will be modified at either the carboxyl terminus or at an internal amino acid. Alternatively, if the therapeutically active region is located at the carboxyl terminus, the therapeutic peptide will be modified at either the amino terminus or at an internal amino acid. Finally, if the therapeutically active region is located at an internal region, the therapeutic peptide will be modified at either the amino terminus or the carboxyl terminus.

"Therapeutic activity" is any activity directed toward healing or curing a biological disorder in a patient. Examples of said therapeutic peptides include pituitary hormones such as vasopressin, oxytocin, melanocyte stimulating hormones, adrenocorticotropic hormones, growth hormones; hypothalamic hormones such as growth hormone releasing factor, corticotropin releasing factor, prolactin releasing

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peptides, gonadotropin releasing hormone and its associated peptides, luteinizing hormone release hormones, thyrotropin releasing hormone, orexin, and somatostatin; thyroid hormones such as calcitonins, calcitonin precursors, and calcitonin gene related peptides; parathyroid hormones and their related proteins; pancreatic hormones such as insulin and insulin-like peptides, glucagon, somatostatin, pancreatic polypeptides, amylin, peptide YY, and neuropeptide Y; digestive hormones such as gastrin, gastrin releasing peptides, gastrin inhibitory peptides, cholecystokinin, secretin, motilin, and vasoactive intestinal peptide: natriuretic peptides such as atrial natriuretic peptides, brain natriuretic peptides, and C-type natriuretic peptides; neurokinins such as neurokinin A. neurokinin B. and substance P; renin related peptides such as renin substrates and inhibitors and angiotensins; endothelins, including big endothelin, endothelin A receptor antagonists, and sarafotoxin peptides; and other peptides such as adrenomedullin peptides, allatostatin peptides, amyloid beta protein fragments, antibiotic and antimicrobial peptides, apoptosis related peptides, bag cell peptides, bombesin, bone Gla protein peptides, CART peptides, chemotactic peptides, cortistatin peptides, fibronectin fragments and fibrin related peptides. FMRF and analog peptides, galanin and related peptides, growth factors and related peptides, Gtherapeutic peptide-binding protein fragments, guanylin and uroguanylin, inhibin peptides, interleukin and interleukin receptor proteins, laminin fragments, leptin fragment peptides, leucokinins, mast cell degranulating peptides, pituitary adenylate cyclase activating polypeptides, pancreastatin, peptide T, polypeptides, virus related peptides, signal transduction reagents, toxins, and miscellaneous peptides such as adjuvant peptide analogs, alpha mating factor, antiarrhythmic peptide, antifreeze polypeptide, anorexigenic peptide, bovine pineal antireproductive peptide, bursin, C3 peptide P16, tumor necrosis factor, cadherin peptide, chromogranin A fragment, contraceptive tetrapeptide, conantokin G, conantokin T, crustacean cardioactive peptide, C-telopeptide, cytochrome b588

peptide, decorsin, delicioius peptide, delta-sleep-inducing peptide, diazempam-binding inhibitor fragment, nitric oxide synthase blocking peptide, OVA peptide, platelet calpain inhibitor (P1), plasminogen activator inhibitor 1, rigin, schizophrenia related peptide, serum thymic factor, sodium potassium Atherapeutic peptidease inhibiro-1, speract, sperm activating peptide, systemin, thrombin receptor agonist, thymic humoral gamma2 factor, thymopentin, thymosin alpha 1, thymus factor, tuftsin, adipokinetic hormone, uremic pentapeptide and other therapeutic peptides.

Taking into account these definitions, the focus of this invention is to modify therapeutic peptides to protect them from peptidase activity *in vivo* and thereby extend the effective therapeutic life of the therapeutic peptide in question as compared to administration of the peptide per se to a patient.

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1. Therapeutic Peptides Used in the Present Invention

Peptide fragments chosen from the determined amino acid sequence of a therapeutic peptide as provided in the attached SEQUENCE LISTING constitute the starting point in the development comprising the present invention. The peptides range from 2 to 50 amino acids in length. The interchangeable terms "peptide fragment" and "peptide moiety" are meant to include both synthetic and naturally occurring amino acid sequences derivable from a naturally occurring amino acid sequence.

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In one embodiment, peptide and peptide fragments are synthesized by conventional means, either by bench-top methods or by automated peptide synthesis machines as discussed in detail below. However, it is also possible to obtain fragments of the peptides by fragmenting the naturally occurring amino acid sequence, using, for example, a proteolytic enzyme. Further, it is possible to obtain the desired fragments of the therapeutic peptide through the use of recombinant DNA technology, as disclosed by Maniatis, T., et al.,

Molecular Biology: A Laboratory Manual, Cold Spring Harbor, New York (1982), which is hereby incorporated by reference. The use of other new modifications to existing methodologies is also contemplated.

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The present invention includes peptides which are derivable from the naturally occuring sequence of the therapeutic peptide. A peptide is said to be "derivable from a naturally occurring amino acid sequence" if it can be obtained by fragmenting a naturally occurring sequence, or if it can be synthesized based upon a knowledge of the sequence of the naturally occurring amino acid sequence or of the genetic material (DNA or RNA) which encodes this sequence. Included within the scope of the present invention are those molecules which are said to be "derivatives" of a peptide. Such a "derivative" has the following characteristics: (1) it shares substantial homology with the therapeutic peptide or a similarly sized fragment of the peptide and (2) it is capable of functioning with the same therapeutic activity as the peptide.

A derivative of a peptide is said to share "substantial homology" with the peptide if the amino acid sequences of the derivative is at least 80%, and more preferably at least 90%, and most preferably at least 95%, the same as that of either the peptide or a fragment of the peptide having the same number of amino acid residues as the derivative.

The derivatives of the present invention include fragments which, in addition to containing a sequence that is substantially homologous to that of a naturally occurring therapeutic peptide may contain one or more additional amino acids at their amino and/or their carboxy termini as discussed in detail below. Thus, the invention pertains to polypeptide fragments of the therapeutic peptide that may contain one or more amino acids that may not be present in a naturally occurring therapeutic peptide sequence provided that such fragments have a therapeutic activity which exceeds that of the therapeutic peptide.

Similarly, the invention includes polypeptide fragments which, although containing a sequence that is substantially homologous to that of a naturally occurring therapeutic peptide, may lack one or more

additional amino acids at their amino and/or their carboxy termini that are naturally found on the therapeutic peptide. Thus, the invention pertains to polypeptide fragments of the therapeutic peptide that may lack one or more amino acids that are normally present in the naturally occurring peptide sequence provided that such polypeptides have a therapeutic activity which exceeds that of the therapeutic peptide.

The invention also encompasses the obvious or trivial variants of the above-described fragments which have inconsequential amino acid substitutions (and thus have amino acid sequences which differ from that of the natural sequence) provided that such variants have an activity which is substantially identical to that of the above-described derivatives. Examples of obvious or trivial substitutions include the substitution of one basic residue for another (i.e. Arg for Lys), the substitution of one hydrophobic residue for another (i.e. Leu for Ile), or the substitution of one aromatic residue for another (i.e. Phe for Tyr), etc.

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As is known in the art, the amino acid residues may be in their protected or unprotected form, using appropriate amino or carboxyl protecting groups as discussed in detail below. The variable length peptides may be in the form of the free amines (on the N-terminus), or acid-addition salts thereof. Common acid addition salts are hydrohalic acid salts, i.e., HBr, HI, or, more preferably, HCl. Useful cations are alkali or alkaline earth metallic cations (i.e., Na, K, Li, Ca, Ba, etc.) or amine cations (i.e., tetraalkylammonium, trialkylammonium, where alkyl can be C_1C_{12}).

Any peptide having a therapeutic activity may be used in this invention. The following list of peptides provides examples of peptides that may be used in this invention, but is not exhaustive and in no way limits the number or type of peptides that may be used in this invention. These therapeutic peptides and fragments produced from these peptides may be modified according to the present invention, and used therapeutically in the body.

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A. Pituitary Hormones (SEQ ID NOS: 1-72)

Adrenocortiocotropic Hormones (ACTH, aka corticotropin)

(SEQ ID NOS: 1-22) - The endocrine functions of the adrenal cortex are regulated by an anterior pituitary hormone, ACTH. ACTH, a 39-amino acid peptide is generated in the corticotrophic cells of the anterior pituitary under the control of corticotropin releasing factor. ACTH is derived by post-translational modification from a 241-amino acid precursor known as pro-opiomelanocortin (POMC).

The biological role of ACTH is to maintain the bulk and the viability of the adrenal cortex and to stimulate the production of adrenal cortex steroids, principally cortisol and costicosterone. The mechanism of action of ACTH involves binding to the ACTH receptor followed by activation of adenylate cyclase, elevation of cyclic AMP (cAMP), and increased protein kinase A (PKA) activity of adrenal cortex tissue. The main effect of these events is to increase the activity of a side chain-cleaving enzyme, which converts cholesterol to pregnenolone. Because of the distribution of enzymes in the various adrenal cortex subdivisions, the principal physiological effect of ACTH is production of the glucocorticosteroids.

Aside from its function controlling adrenal cortical activity, ACTH appears to have diverse biological roles including modulation of endocrine and exocrine glands, temperature regulation and influences on nerve regeneration and development. In addition, ACTH and its fragments affect motivation, learning, and behavior. The use of ACTH as a therapeutic agent may thus help the control of these functions. ACTH release from the anterior pituitary is mediated by corticotropin releasing factor (CRF).

Growth Hormone Peptides (SEQ ID NOS: 23-24, 45) – Human placental lactogen (hPL), growth hormones, and prolactin (Prl) comprise the growth hormone family. All have about 200 amino acids, 2 disulfide bonds, and no glycosylation. Although each has special receptors and

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unique characteristics to their activity, they all possess growth-promoting and lactogenic activity. Mature GH (22,000 daltons) is synthesized in acidophilic pituitary somatotropes as a single polypeptide chain. Because of alternate RNA splicing, a small amount of a somewhat smaller molecular form is also secreted.

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There are a number of genetic deficiencies associated with GH. GH-deficient dwarfs lack the ability to synthesize or secrete GH, and these short-statured individuals respond well to GH therapy. Pygmies lack the IGF-1 response to GH but not its metabolic effects; thus in pygmies the deficiency is post-receptor in nature. Finally, Laron dwarfs have normal or excess plasma GH, but lack liver GH receptors and have low levels of circulating IGF-1. The defect in these individuals is clearly related to an inability to respond to GH by the production of IGF-1. The production of excessive amounts of GH before epiphyseal closure of the long bones leads to gigantism, and when GH becomes excessive after epiphyseal closure, acral bone growth leads to the characteristic features of acromegaly. Using GH as a therapeutic agent would aid in treating these disorders, and potentially stimulate growth in other cases of short stature with low or normal GH levels.

Melanocyte Stimulating Hormones (MSH) (SEQ ID NOS: 25-39) - Melanocyte stimulating hormone (MSH) is generated in the intermediary pituitary under the control of dopamine. MSH may have important physiological roles in the control of vertebrate pigment cell melanogenesis, neural functioning related to learning and behavior, and fetal development. See Sawyer, T.K. et al., Proc. Nat. Acad. Sci USA, 79, 1751 (1982).

Oxytocin (SEQ ID NOS: 40-44) - Oxytocin is involved in the enhancement of lactation, contraction of the uterus, and relaxation of the pelvis prior to childbirth. Oxytocin secretion in nursing women is stimulated by direct neural feedback obtained by stimulation of the nipple during suckling. Its physiological effects include the contraction of mammary gland myoepithelial cells, which induces the ejection of milk

from mammary glands, and the stimulation of uterine smooth muscle contraction leading to childbirth. Oxytocin causes myoepithelial cells surrounding secretory acini of mammary glands to contract, pushing milk through ducts. In addition, it stimulates the release of prolactin, and prolactin is trophic on the breast and stimulates acinar formation of milk. A conjugated oxytocin could thus be used to aid lactation and help relax the pelvis prior to birth. It could also be used to prevent post partum uterine hemorrage.

Vasopressin (ADH) (SEQ ID NOS: 46-72) – Vasopressin is also known as antidiuretic hormone (ADH), because it is the main regulator of body fluid osmolarity, causing antidiuresis and increase in blood pressure. Vasopressin binds plasma membrane receptors and acts through G-proteins to activate the cyclic AMP/protein kinase A (cAMP/PKA) regulatory system. The secretion of vasopressin is regulated in the hypothalamus by osmoreceptors, which sense water concentration and stimulate increased vasopressin secretion when plasma osmolarity increases. The secreted vasopressin increases the reabsorption rate of water in kidney tubule cells, causing the excretion of urine that is concentrated in Na⁺ and thus yielding a net drop in osmolarity of body fluids. Vasopressin deficiency leads to watery urine and polydipsia, a condition known as diabetes insipidus. Using conjugated vasopressin or vasopressin fragments would thus prevent these disorders and allow the regular maintenance of the body's osmolarity.

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B. Hypothalamic Hormones (Releasing factors) Corticotropin Releasing Factor (CRF) & related peptides (SEQ ID NOS: 73-102) – Corticotrophin-releasing factor (CRF), a 41 amino acid peptide, plays a significant role in coordinating the overall response to stress through actions both in the brain and the periphery. In the brain, CRF is produced and secreted primarily from parvocellular neurons of the paraventricular hypothalamic nucleus. From there, the

CRF-containing neurons project to the portal capillary zone of the median eminence and act to stimulate the secretion of adrenocorticotrophic hormone (ACTH), beta-endorphin, and other proopiomelanocortin (POMC)-derived peptides from the pituitary gland.

The subsequent ACTH-induced release of adrenal glucocorticoids represents the final stage in the hypothalamic-pituitary-adrenal axis (HPA), which mediates the endocrine response to stress. Besides its neuroendocrine role, CRF also functions as a neurotransmitter and neuromodulator to elicit a wide spectrum of autonomic, behavioral and immune effects to physiological, pharmacological, and pathological stimuli.

Clinical studies indicated that CRF hypersecretion is associated with various diseases, such as major depression, anxiety-related illness, eating disorder, as well as inflammatory disorder. Low levels of CRF have been found in Alzheimer's disease, dementias, obesity, and many endocrine diseases. Therefore, the use of CRF as a therapeutic agent to counter the effects associated with high levels or low levels of CRF will provide a basis for the treatment of diseases that are associated with abnormal CRF levels. Several peptide antagonists and nonpeptide antagonists have been discovered and widely studied, including a-helical CRF(9-41), Astressin, D-PheCRF(12-41) (peptide antagonist) and CP-154526 (nonpeptide antagonist). These CRF antagonists may provide a novel agent for treatment of depression, anxiety and other CRF related illnesses. Conjugated CRF peptides could thus be used to maintain adrenal health and viability during long term steroid use or as anti-inflamatory agents.

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Gonadotropin Releasing Hormone Associated peptides
(GAP) (SEQ ID NOS: 103-110) - GAP is contained in the precursor molecule to gonadotropin-releasing hormone (GnRH). GAP has prolactin inhibiting properties. Gn-RH is a hormone secreted by the hypothalamus that stimulate the release of gonadotrophic hormones follicle stimulating hormone (FSH) and luteinizing hormone (LH). Low

levels of circulating sex hormone reduce feedback inhibition on GnRH synthesis, leading to elevated levels of FSH and LH. The latter peptide hormones bind to gonadal tissue, resulting in sex hormone production via cyclic AMP (cAMP) and protein kinase A (PKA) mediated pathways. A conjugated GnRM could be used to aid fertility, or as a contraceptive

in either males or females. This agent would have use in animals as well as humans.

Growth Hormone Releasing Factor (GRF) (SEQ ID NOS: 111-134)- GRF is a hypothalamic peptide that plays a critical role in controlling the synthesis and secretion of growth hormone in the anterior pituitary. Some structurally unrelated short peptides have also been

reported to elicit growth hormone secretion by a different mechanism.

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Under the influence of GRF, growth hormone is released into the systemic circulation, causing the target tissue to secrete IGF-1. Growth hormone also has other more direct metabolic effects; it is both hyperglycemic and lipolytic. The principal source of systemic IGF-1 is the liver, although most other tissues secrete and contribute to systemic IGF-1. Liver IGF-1 is considered to be the principal regulator of tissue growth. In particular, the IGF-1 secreted by the liver is believed to synchronize growth throughout the body, resulting in a homeostatic balance of tissue size and mass. IGF-1 secreted by peripheral tissues is generally considered to be autocrine or paracrine in its biological action. The use of a conjugated GRF as a therapeutic agent to increase GH release, would then help treat disorders involving growth functions regulated by GRF.

Lutenizing Hormone Release Hormones (LH-RH) (SEQ ID NOS: 135-161) – Luteinizing hormone releasing hormone is the key mediator in the neuroregulation of the secretion of gonadotropins, luteinizing hormone (LH) and follicle stimulating hormone (FSH). LH-RH can modify sexual behavior by regulating plasma gonadotropin and sex steroid levels. See Vale, W.W. et al., Peptides, Structure and Biological Function, Proceedings of the Sixth American Peptide Symposium,

Gross, E. and Meienhofer, M., eds., 781 (1979). A conjugated LH-RH agent could be used to stimulate ovulation in humans or animals as an aid to fertility.

Orexins (SEQ ID NOS: 162-164) – Orexins are a family of neuropeptides from the hypothalamus that have been recently discovered and characterized. Orexins stimulate appetite and food consumption. Their genes are expressed bilaterally and symmetrically in the lateral hypothalamus, which was earlier determined to be the "feeding center" of the hypothalamus. In contrast, the so-called satiety center is expressed in the ventromedial hypothalamus and is dominated by the leptin-regulated neuropeptide network.

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Prolactin Releasing Peptides (SEQ ID NOS: 65-170)— Prolactin is produced by acidophilic pituitary lactotropes. Prolactin releasing peptides act on lactotrope to release prolactin. PRL initiates and maintains lactation in mammals, but normally only in mammary tissue that has been primed with estrogenic sex hormones. A conjugated PRP could be used to increase lactation in humans or animals.

Somatostatin (SEQ ID NOS: 171-201) – Also known as Growth Hormone Release Inhibiting Factors (GIF), somatostatin is a 14 amino acid peptide is secreted by both the hypothatamus and by d cells of the pancreas (its pancreatic version is discussed below). Somatostatin has been reported to modulate physiological functions at various sites including pituitary, pancreas, gut and brain. It inhibits the release of growth hormone, insulin, and glucagon. It has many biological roles, including: inhibition of basal and stimulated hormone secretion from endocrine and exocrine cells, an effect on locomotor activity and cognitive function, and possible therapeutic value in small cell lung cancer. See Reubi, J. C. et al, Endocrinology, 110, 1049 (1982). A conjugated somatostatin could be used to treat giantism in children or acromegaly in the adult.

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Thyrotropin Releasing Hormone (THR) and Analogs (SEQ ID

NOS: 202-214) – THR stimulates the production of thyroid stimulating hormone (TSH, also known as thyrotropin) and prolactin secretion. In adults, TSH is responsible for up-regulating general protein synthesis and inducing a state of positive nitrogen balance. In the embryo, it is necessary for normal development. Hypothyroidism in the embryo is responsible for cretinism, which is characterized by multiple congenital defects and mental retardation. A conjugated THR could then be used as a therapeutic agent in the treatment of these disorders. It could also be used to treat pituitary causes of thyroid insufficiency or in the diagnosis of human tumors of the thyroid.

C. Thyroid Hormones

Calcitonins (CT) & Caltitonins Precursor Peptides (SEQ ID

NOS: 215-224) - Calcitonin (CT) is a 32-amino acid peptide secreted by C cells of the thyroid gland. Calcitonin is employed therapeutically to relieve the symptoms of osteoporosis, although details of its mechanism of action remain unclear. However, it has been observed that CT induces the synthesis of parathyroid hormone (PTH) in isolated cells, which leads *in vivo* to increased plasma Ca²⁺ levels. In addition, CT has been shown to reduce the synthesis of osteoporin (Opn), a protein made by osteoclasts and responsible for attaching osteoclasts to bone. Thus, using conjugated CT as a therapeutic peptide would elevate plasma Ca²⁺ via PTH induction and reduce bone reabsorption by decreasing osteoclast binding to bone.

Calcitonins Gene Related Peptide (CGRP) (SEQ ID NOS: 225-

<u>253</u>)— CGRP is a 37 amino acid peptide that results from alternative splicing of calcitonin gene transcripts. It exists in at least two forms: alpha-CGRP (or CGRP-I) and beta-CGRP (or CGRP-II). CGRP has considerable homology with amylin and adrenomedullin, and is widely distributed both centrally and peripherally in organs including the skin,

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the heart, the pancreas, the lungs, and the kidneys. CGRP has many biological roles, affecting the nervous and cardiovascular systems, inflammation and metabolism.

5 D. Parathyroid Hormones and Related Proteins Parathyroid Hormones (PTH) (SEQ ID NOS: 254-293) -

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Parathyroid hormone (PTH) is synthesized and secreted by chief cells of the parathyroid in response to systemic Ca²⁺ levels. It plays a major role in the modulation of serum calcium concentration and thereby affect the physiology of mineral and bone metabolism. The Ca²⁺ receptor of the parathyroid gland responds to Ca²⁺ by increasing intracellular levels of PKC, Ca²⁺ and IP₃; this stage is followed, after a period of protein synthesis, by PTH secretion. The synthesis and secretion of PTH in chief cells is constitutive, but Ca²⁺ regulates the level of PTH in chief cells (and thus its secretion) by increasing the rate of PTH proteolysis when plasma Ca²⁺ levels rise and by decreasing the proteolysis of PTH when Ca²⁺ levels fall. The role of PTH is to regulate Ca²⁺ concentration in extracellular fluids. The feedback loop that regulates PTH secretion therefore involves the parathyroids, Ca²⁺, and the target tissues described below.

PTH acts by binding to cAMP-coupled plasma membrane receptors, initiating a cascade of reactions that culminates in the biological response. The body's response to PTH is complex but is aimed in all tissues at increasing Ca²⁺ levels in extracellular fluids. PTH induces the dissolution of bone by stimulating osteoclast activity, which leads to elevated plasma Ca²⁺ and phosphate. In the kidney, PTH reduces renal Ca²⁺ clearance by stimulating its reabsorption; at the same time, PTH reduces the reabsorption of phosphate and thereby increases its clearance. Finally, PTH acts on the liver, kidney, and intestine to stimulate the production of the steroid hormone 1,25-dihydroxycholecalciferol (calcitriol), which is responsible for Ca²⁺ absorption in the intestine. A conjugated PTH could be used to regulate

calcium homeostasis in patients with parathyroid hormone deficiency states. Inhibitor analogues could be used to block PTH action in renal failure or other patients with excessive PTH levels.

Parathyroid Hormone Related Proteins (PTHrP) (SEQ ID NOS:

294-309) - Parathyroid hormone-related protein (PTHrP) has received 5 attention as a physiological regulator attenuating chondrocytic differentiation and preventing apoptotic cell death. PTHrP was initially identified as a tumor-derived, secretory protein with structural similarity to parathyroid hormone (PTH), the major regulator of calcium homeostasis. PTH and PTHrP bind to a common G protein-coupled cell 10 surface receptor (PTH/PTHrP or PTH-1 receptor) that recognizes the Nterminal (1-34) region of these peptides. Hence, when tumor-derived PTHrP enters the circulation, it activates receptors in classic PTH target organs such as bone and kidney and elicits PTH-like bioactivity. By promoting bone resorption and inhibiting calcium excretion, circulating 15 PTHrP gives rise to the common paraneoplastic syndrome of malignancy-associated humoral hypercalcemia.

Although initially discovered in tumors, PTHrP was subsequently shown to be expressed in a remarkable variety of normal tissues including the fetal and adult skeleton, where acting in concert with its amino terminal PTH-1 receptor, it serves to regulate cellular growth and differentiation. The anabolic effects of intermittent PTH administration on bone and its therapeutic potential in osteoporosis have been extensively explored. With the recognition that PTHrP is the endogenous ligand for the PTH/PTHrP receptor in osteoblasts, its use as an anabolic agent has also been investigated. Modified PTHrP peptides could be used for similar indications as PTH.

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E. Pancreatic Hormones - The principal role of the pancreatic
30 hormones is the regulation of whole-body energy metabolism, principally

by regulating the concentration and activity of numerous enzymes involved in catabolism and anabolism of the major cell energy supplies.

Amylin (SEQ ID NOS: 310-335)- Pancreatic beta-cell hormone amylin is a 37-amino-acid peptide related to CGRP and calcitonin. It is co-secreted with insulin from pancreatic beta-cells. Amylin is deficient with type 1 diabetes mellitus. Amylin appears to work with insulin to regulate plasma glucose concentrations in the bloodstream, suppressing the postherapeutic peptiderandial secretion of glucagon and restraining the rate of gastric emptying. People with diabetes have a deficiency in the secretion of amylin that parallels the deficiency in insulin secretion, resulting in an excessive inflow of glucose into the bloodstream during the postherapeutic peptiderandial period.

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While insulin replacement therapy is a cornerstone of diabetes treatment, replacement of the function of both amylin and insulin may allow a more complete restoration of the normal physiology of glucose control. Type 2 diabetes is characterized by islet amyloid deposits, which are primarily composed of the amyloidogenic human form of islet amyloid polypeptide. A conjugated amylin could be used in the management of diabetes to limit post prandial hyperglysemia.

Glucagon (SEQ ID NOS: 336-376) - Glucagon is a 29-amino acid hormone synthesized by the a cells of the islets of Langerhans as a very much larger proglucagon molecule. Like insulin, glucagon lacks a plasma carrier protein, and like insulin its circulating half life is also about 5 minutes. As a consequence of the latter trait, the principal effect of glucagon is on the liver, which is the first tissue perfused by blood containing pancreatic secretions. Glucagon binds to plasma membrane receptors and is coupled through G-proteins to adenylate cyclase. The resultant increases in cAMP and PKA reverse all of the effects described above that insulin has on liver. The increases also lead to a marked elevation of circulating glucose, with the glucose being derived from liver gluconeogenesis and liver glycogenolysis. A conjugated glucagon

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construct could be used to manage brittle diabetes with recurrent hypoglycemia or to prevent or treat iatrogenic hypoglycemia.

Insulin and Insulin-Like peptides (SEQ ID NOS: 377-382) -

The earliest of these hormones recognized was insulin, a disulfide bonded dipeptide of 21 and 30 amino acids produced by the pancreas, whose major function is to counter the concerted action of a number of hyperglycemia-generating hormones and to maintain low blood glucose levels. Insulin is a member of a family of structurally and functionally similar molecules that include IGF-1, IGF-2, and relaxin. The tertiary structure of all 4 molecules is similar, and all have growth-promoting activities, but the dominant role of insulin is metabolic while the dominant roles of the IGFs and relaxin are in the regulation of cell growth and differentiation.

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Insulin is synthesized as a preprohormone in the b cells of the islets of Langerhans. Its signal peptide is removed in the cisternae of the endoplasmic reticulum and it is packaged into secretory vesicles in the Golgi, folded to its native structure, and locked in this conformation by the formation of 2 disulfide bonds. Specific protease activity cleaves the center third of the molecule, which dissociates as C peptide, leaving the amino terminal B peptide disulfide bonded to the carboxy terminal A peptide.

Insulin generates its intracellular effects by binding to a plasma membrane receptor, which is the same in all cells. The receptor is a disulfide-bonded glycoprotein. One function of insulin (aside from its role in signal transduction) is to increase glucose transport in extrahepatic tissue is by increasing the number of glucose transport molecules in the plasma membrane. Glucose transporters are in a continuous state of turnover. Increases in the plasma membrane content of transporters stem from an increase in the rate of recruitment of new transporters into the plasma membrane, deriving from a special pool of preformed transporters localized in the cytoplasm.

In addition to its role in regulating glucose metabolism (and its therapeutic use in treating diabetes), insulin stimulates lipogenesis, diminishes lipolysis, and increases amino acid transport into cells. Insulin also modulates transcription, altering the cell content of numerous mRNAs. It stimulates growth, DNA synthesis, and cell replication, effects that it holds in common with the IGFs and relaxin. A conjugated insulin could thus be used to manage diabetes.

NeuroPeptide Y (SEQ ID NOS: 383-389) – Neuropeptide Y (NPY), a peptide with 36 amino acid residues, is one of the most abundant neuropeptides in both the peripheral and the central nervous systems. It belongs to the pancreatic polypeptide family of peptides. Like its relatives, peptide YY (PYY) and pancreatic polypeptide (PP), NPY is bent into hairpin configuration that is important in bringing the free ends of the molecule together for binding to the receptors.

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NPY exerts a wide range of effects in the central nervous system (CNS) and the periphery. Its CNS actions include major effects on feeding and energy expenditure, and alterations in heart rate, blood pressure, arousal and mood. In the periphery, NPY causes vasoconstriction and hypertension; it is also found in the gastrointestinal and urogenital tract, implicating its functions by action upon gastrointestinal and renal targets. In recent studies, hypothalamic NPY has been found to play a fundamental role in developing the features of obesity, it is a major transducer in the pathways signalling body fat to the hypothalamus, and in regulating body fat content. Leptin, an obese gene product, has been found to decrease NPY gene expression in obese (ob/ob) mice. Insulin and corticosteroids are also involved in the regulation of hypothalamic NPY synthesis, with insulin decreasing and corticosteroids increasing NPY expression. A conjugated NPY could be used to treat obesity and MODM (Type II DM) in obese patients.

Pancreatic Polypeptides (PP) (SEQ ID NOS: 390-396) Pancreatic polypeptide (PP) is a 36-amino acid hormone produced by F
cells within the pancreatic islets and the exocrine pancreas. It is a

member of the PP fold family of regulatory peptides, and increases glycogenolysis and regulates gastrointestinal activity. A conjugated pancreatic polypeptide could thus be used to alter absorption and metabolism of foods.

Peptide YY (SEQ ID NOS: 397-400) - PYY is a thirty six amino acid long peptide, first isolated from porcine intestinal tissue and mainly localized in intestinal endocrine cells. It has many biological activities, including a range of activities within the digestive system and potent inhibition of intestinal electrolyte and fluid secretion.

Somatostatin (SEQ ID NOS: 171-201) — The somatostatin secreted by d cells of the pancreas is a 14-amino acid peptide identical to somatostatin secreted by the hypothalamus. In neural tissue somatostatin inhibits GH secretion and thus has systemic effects. In the pancreas, somatostatin acts a paracrine inhibitor of other pancreatic hormones and thus also has systemic effects. It has been speculated that somatostatin secretion responds principally to blood glucose levels, increasing as blood glucose levels rise and thus leading to down-regulation of glucagon secretion. A conjugated somatostatin could then be used to aid in the management of diabetes.

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F. Digestive Hormones

Cholecystokinin (CCK) & related peptides (SEQ ID NOS: 401-

<u>416)</u> – CCK is a polypeptide of 33 amino acids originally isolated from pig small intestine that stimulates gallbladder contraction and bile flow and increases secretion of digestive enzymes from pancreas. It exists in multiple forms, including CCK-4 and CCK-8, with the octapeptide representing the dominant molecular species showing the greatest activity. It belongs to the CCK/gastrin peptide family and is distributed centrally in the nervous system and peripherally in the gastrointestinal system. It has many biological roles, including stimulation of pancreatic secretion, gall bladder contraction and intestinal mobility in the GI tract

as well as the possible mediation of satiety and painful stimuli. A conjugated CCK could be used in diagnostic studies of the gall bladder or in chronic cholecystisis.

Gastrin Releasing Peptide (GRP) (SEQ ID NOS: 417-429) -

GRP is a 27-amino acid peptide originally isolated from porcine non-antral gastric tissue, and is the homolog of the frog skin peptide named bombesin growth. It is widely distributed both centrally and peripherally in tissues including brain, lung and gastrointestinal tract. It regulates a variety of cell physiological processes including secretion, smooth muscle contraction, neurotransmission and cell growth. A conjugated GRP could be used in the treatment of adynamic ileus or constipation in the elderly.

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Gastrin & related peptides (SEQ ID NOS: 417-429) - Gastrin is a polypeptide of 17 amino acids produced by stomach antrum, which stimulates acid and pepsin secretion. Gastrin also stimulates pancreatic secretions. Multiple active products are generated from the gastrin precursor, and there are multiple control points in gastrin biosynthesis. Biosynthetic precursors and intermediates (progastrin and Gly-gastrins) are putative growth factors; their products, the amidated gastrins. regulate epithelial cell proliferation, the differentiation of acid-producing parietal cells and histamine-secreting enterochromaffin-like (ECL) cells. and the expression of genes associated with histamine synthesis and storage in ECL cells, as well as acutely stimulating acid secretion. Gastrin also stimulates the production of members of the epidermal growth factor (EGF) family, which in turn inhibit parietal cell function but stimulate the growth of surface epithelial cells. Plasma gastrin concentrations are elevated in subjects with Helicobacter pylori, who are known to have increased risk of duodenal ulcer disease and gastric cancer. The use of gastrin or gastrin antagonists as a therapeutic agent may therefore contribute to treating major upper gastrointestinal tract disease.

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Gastrin Inhibitory peptides (SEQ ID NOS: 417-429) – Gastrin inhibitory peptide is a polypeptide of 43 amino acids that inhibits secretion of gastrin. A conjugated GIP could be used to treat severe peptic ulcer disease.

Motilin (SEQ ID NOS: 430-433) – Motilin is a polypeptide of 22 amino acids that controls gastrointestinal muscles. Motilin-producing cells are distributed in the duodenum, upper jejunum, and colorectal adenocarinomas and in midgut carcinoids. Motilin stimulates gut motility.

Secretin (SEQ ID NOS: 434-441) – Secretin is a polypeptide of 27 amino acids secreted from duodenum at pH values below 4.5, stimulates pancreatic acinar cells to release bicarbonate and H₂O. Secretin is a neurotransmitter (a chemical messenger) in the neuropeptide group. It is one of the hormones that controls digestion (gastrin and cholecystokinin are the others). It is a polypeptide composed of 27 amino acids and is secreted by cells in the digestive system when the stomach empties. Secretin stimulates the pancreas to emit digestive fluids that are rich in bicarbonate which neutralizes the acidity of the intestines, the stomach to produce pepsin (an enzyme that aids digestion of protein), and the liver to produce bile.

Secretin may be useful in treating autism. In one study, children with autistic spectrum disorders underwent upper gastrointestinal endoscopy and intravenous administration of secretin to stimulate pancreaticobiliary secretion. All three had an increased pancreaticobiliary secretory response when compared with nonautistic patients (7.5 to 10 mL/min versus 1 to 2 mL/min). Within 5 weeks of the secretin infusion, a significant amelioration of the children's gastrointestinal symptoms was observed, as was a dramatic improvement in their behavior, manifested by improved eye contact, alertness, and expansion of expressive language. These clinical observations suggest an association between gastrointestinal and brain function in patients with autistic behavior.

Vasoactive Intestinal Peptide (VIP) and related peptides

WO 00/69900 PCT/US00/13576

(SEQ ID NOS: 442-464) –VIP is a polypeptide of 28 residues produced by hypothalamus and GI tract. It relaxes the GI, inhibits acid and pepsin secretion, acts as a neurotransmitter in peripheral autonomic nervous system, and increases secretion of H₂O and electrolytes from pancreas

and gut. It was originally discovered in lung and intestine and is also found in tissues including brain, liver, pancreas, smooth muscle and lymphocytes. It is structurally related to a family of peptides which

include PACAP, PHI, secretin and glucagon. It has a diverse range of

biological actions including vasodilation, electrolyte secretion, modulation of immune function and neurotransmission. A conjugated VIP may be useful in the treatment of achlorhydria, ischemic colitis and irritable bowel syndrome (IBS).

- 15 **G.** Natriuretic Peptides There are three members in the natriuretic peptide hormone family, atrial natriuretic peptide (ANP), B-type natriuretic peptide (BNP, brain natriuretic peptide), and C-type natriuretic peptide (CNP), that are involved in the regulation of blood pressure and fluid homeostasis.
- Atrial-Natriuretic Peptides (ANP) (SEQ ID NOS: 465-507)-20 ANP is a 28-amino acid peptide hormone containing a disulfide bond. It exerts natriuretic, diuretic, and vasorelaxant effects and play an important role in the body's blood volume and blood pressure homeostasis. See Smith, F.G. et al., J. Dev. Physiol. 12, 55 (1989). The mechanisms controlling ANP release have been the subject of intense 25 research, and are now fairly well understood. The major determinant of ANP secretion is myocyte stretch. Although much less is known about the factors regulating BNP release from the heart, myocyte stretch has also been reported to stimulate BNP release from both atria and ventricles. However, whether wall stretch acts directly or via factors such 30 as endothelin-1, nitric oxide, or angiotensin II liberated in response to distension has not been established. Recent studies show that by

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stimulating endothelin type A receptors endothelin plays an important physiological role as a mediator of acute-volume load-induced ANP secretion from atrial myocytes in conscious animals. In fact, endogenous paracrine/autocrine factors liberated in response to atrial wall stretch rather than direct stretch appears to be responsible for activation of ANP secretion in response to volume load, as evidenced by almost complete blockade of ANP secretion during combined inhibition of endothelin type A/B and angiotensin II receptors. Furthermore, under certain experimental conditions angiotensin II and nitric oxide may also exert a significant modulatory effect on stretch-activated ANP secretion. The molecular mechanisms by which endothelin-1, angiotensin II, and nitric oxide synergistically regulate stretch-activated ANP release are yet unclear. Abstract Volume 75 Issue 11/12 (1997) pp 876-885, Journal of Molecular Medicine. A conjugated would be useful in the management of malignant hypertension or severe hypertension and renal failure.

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Brain Natriuretic Peptides (BNP) (SEQ ID NOS: 507-516) Brain natriuretic peptide (BNP), a member of the natriuretic peptide
family, is produced and released from cardiac ventricles. BNP regulates
the body fluid volume, blood pressure, and vascular tones through the Atype guanylate cyclase-coupled receptor. The BNP plays a role in
electrolyte-fluid homeostasis such as atrial natriuretic peptides (ANP). A
conjugated BNP could be useful in the management of heart failure.

C-Type Natriuretic Peptides (CNP) (SEQ ID NOS: 517-524) - C-type natriuretic peptide (CNP), the third member of the natriuretic peptide family, is produced in vascular endothelial cells (ECs) and acts as an endothelium-derived relaxing peptide. Although atrial and brain natriuretic peptides are well known to be involved in the regulation of cardiovascular and endocrine functions as circulating hormones, the roles of the C-type natriuretic peptide (CNP) remain unknown.

CNP is found principally in the central nervous system and vascular endothelial cells while ANP and BNP are cardiac hormones.

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ANP is synthesized mainly in the atria of the normal adult heart, while BNP is produced by both the atria and ventricles.

H. Tachykinins (SEQ ID NOS: 525-627) – A family of peptides,
including neurokinin A and substance P, that share a common C
terminal sequence (F-X-GLM-NH2) which is required for full biological activity including Neurokinin A, B, and Substance P.

Neurokinin A – Neurokinin A is a decapeptide, previously known as substance K. It is is a member of the tachykinin family of neuropeptides which includes substance P and neurokinin B. It exhibits a variety of activities related to smooth muscle contraction, pain transmission, bronchoconstriction, vasodilation and modulation of the immune system.

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Neuromedin- Neuromedins, smooth-muscle-stimulating peptides, are commonly divided into four groups: bombesin-like, kassinin-like, neurotensin-like and neuromedins U. These neuropeptides and their receptors are localized to all components of the HPA hypophyseal pituitary axis, the only exemption seems to be neurokinin B, which is not detected in the adenohypophysis. Neuromedins exert a manifold effect on HPA axis, and their action on the adrenal suggests their involvement in the regulation of growth, structure and function of the adrenal cortex. Neuromedins may exert both direct and indirect effects on the adrenal cortex. Direct effect is proven by the stimulation of mineralo- and glucocorticoid therapeutic peptides by isolated or cultured adrenocortical cells and by mobilisation of intracellular [Ca2+]. Indirect effects, on the other hand, may be mediated by ACTH, arginine-vasopressin, angiotensin II, catecholamines or by other regulatory substances of medullary origin.

<u>Substance P and Related Peptides</u> – Substance P is an eleven amino acid peptide, first isolated from brain and intestine. It has been proposed as a neuromodulator involved in pain transmission in the spinal cord. It also affects contraction of smooth muscle, reduction of

blood pressure, stimulation of secretory tissue, and release of histimine from mast cells.

I. **Renin Related Peptides**

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Angiotensins (SEQ ID NOS: 628-677)— Angiotensin is a 10 amino acid peptide derived from enzymatic cleavage of a2-globin by the kidney enzyme renin. The C-terminal 2 amino acids are then released to yield angiotensin I, which is responsible for essential hypertension through stimulated synthesis and release of aldosterone from adrenal cells. It is a multifunctional hormone regulating blood pressure, plasma volume, neuronal function thirst, and water intake.

Angiotensin II is an octapeptide derived from angiotensin I by angiotensin converting enzyme, and is widely distributed both centrally and peripherally in organs such as the heart, the kidneys, and the liver. Angiotensin IV is the terminal hexapeptide fragment of angiotensin II formed metabolically by proteolytic cleavage from either angiotensin I or angiotensin II. It plays a role in vascular control, cardiac growth, renal blood flow and memory function.

Angiotensin II is the key peptide hormone that regulates vascular smooth muscle tone, blood pressure, free water intake and sodium retension. It controls vascular homeostasis compensating for loss of intravascular volume by stimulating increased vasospastic tone, increase sodium retention and increased free water intake.

Renin substrates and Inhibitors (SEQ ID NOS: 678-684) -Renin is a very specific aspartic protease, which is synthesized and released by differentiated smooth muscle cells in the vasculature of the kidney called granular epithelial cells. Renin is specific for its substrate, angiotensinogen, which it cleaves specifically at the Leu¹⁰-Val¹¹ bond to form the decapeptide, angiotensin I (AI). The renin-angiotensin system is involved in the control of fluid and mineral balance throughout the 30 vertebrates. Renin can be found in mammals, birds, reptiles, amphibians, bony fishes, cartilaginous fishes, and agnathans. Specific

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renin inhibitors can also be designed, with therapeutic applications for treatment of for example hypertension and congestive heart failure (Blundell et al., 1987).

J. Endothelins and Related Peptides (SEQ ID NOS: 685-744) - The endothelin peptide family consists of the 21 amino acid isoforms endothelin-1, endothelin-2, endothelin-3, sarafotoxin (a snake venom) and scorpion toxin.

Endothelins (ET) and Big Endothelins – Endothelins are found
 on endothelial cells in a wide variety of organ systems. Examples of pathologies and physiological processes associated with changes in endothelin levels and synthetics include: atherosclerosis and hypertension, coronary vasospasm, acute renal failure, changes in intracellular Ca2⁺ levels, and effects on the renin-angiotensin system.
 Endothelins are released in reponse to variations in angiotensin II, vasopressin, and cytokines (e.g. TGF-βm TNF-α, IL-β-) levels as well as other physiological events including increase blood flow.

The endothelin family of peptides consists of highly potent endogenous vasoconstrictor agents first isolated from endothelial cell supernatant. They regulate blood flow to organs by exporting a vasoconstrictive effect on arteries. Endothelins are derived from bigendothelin, which is cleaved by a unique membrane-bound metalloprotease, endothelin-converting enzyme, into the 21-amino-acid bioactive forms (ET-1, ET-2 and ET-3).

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Of the 3 isoforms (ET-1, ET-2, ET-3), endothelin-1 is the major isoform and plays an important role for regulation of vascular function. Endogenous endothelin peptides and their receptors are differentially distributed throughout the many smooth muscle tissues including blood vessels, uterus, bladder and intestine. Through this widespread distribution and localization, they exert biological functions in regulating vascular tone and causing mitogenesis. ETs and their receptor subtypes are also present in various endocrine organs. It appears to act as a

modulator of secretion of prolactin, gonadotropins GH and TSH. Endothelin may also be the disease marker or an etiologic factor in ischemic heart disease, atherosclerosis, congestive heart failure, renal failure, systemic hypertension, pulmonary hypertension, cerebral vasospasm.

Exogenously administered endothelin-1 has been demonstrated to increase peripheral resistance and blood pressure in a dose-dependent manner. However, during the first minutes of intravenous administration endothelins also decrease peripheral resistance and blood pressure, presumably due to the release of vasodilatory compounds such as nitric oxide, prostacyclin, and atrial natriuretic peptide.

<u>ET(A) Receptor Antagonists</u> – Endothelin receptors exist as two types: A (ET-A) and B (ET-B1 and ET-B2). ET-A receptors are responsible for while ET-B1 and ET-B2 mediate vasorelaxation and vasoconstriction respectively.

<u>Sarafotoxin peptides</u> – As already described, endothelin (ET) peptides are potent growth factors binding to G protein-coupled receptors. Sarafotoxins (S6) isolated from *Atractaspis engaddensis* are highly homologous to endothelins. Sarafotoxin peptides have marked vasoconstrictive activity and are responsible for the ischemic limb loss that follows snake or scorpion bites. They could be used therapeutically as a peptidase stabalized peptide as a vasopressive agent in shock and sepsis.

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K. Opioid Peptides (SEQ ID NOS: 745-927) - Opioids are a large class of drugs, used clinically as painkillers, that include both plant-derived and synthetic alkaloids and peptides found endogenously in the mammalian brain. While the plant-derived alkaloids have been known and used for thousands of years, the endogenous opioid peptides were discovered only in the mid-1970s.

Opioids include casomorphin peptides, demorphins, endorphins, enkephalins, deltorphins, dynorphins, and analogs and derivatives of these.

Casomorphin Peptides – Casomorphin peptides are novel opioid peptides derived from casein(§-casomorphins). Beta casomorphins are the more exstensive studied opioid peptides arising from food proteins (beta-caseins). They were originally isolated from bovine beta-casein, the same sequences occur in ovine and buffalo beta-caseins.

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<u>Dermorphins</u> – Demorphin is a is a seven amino acid peptide, originally isolated from *Phylomedusa sauvagei* frog skin. It is a ligand which binds with high affinity to the μ opioid receptor, and has many biological roles including analgesia, endocrine modulation, immunomodulation, increased K⁺ conductance and inhibition of action potentials.

Dynorphin/New-Endorphin Precursor Related Peptides -Dynorphins are a class of endogenous opioids that exist in multiple forms in the central nervous system. Dynorphins are derived from the precursor prodynorphin (proenkephalin B). Dynorphin, also known as Dynorphin A1-17, is a well known opioid which has the sequence Tyr-Gly-Gly-Phe-Leu⁵-Arg-Arg-Ile-Arg-Pro¹⁰-Lys-Leu-Lys-Trp-Asp¹⁵-Asn-Gln. SEQ ID NO:1. A number of derivatives and analogs of dynorphin are known including Dyn A1-13, SEQ ID NO: 2 Dyn A2-13, SEQ ID NO:3, Dvn A1-12, Dvn A2-12 and Dvn A2-17 as well as amide analogs such as those mentioned in U.S. Patent 4,462,941 of Lee et al., N-terminus truncated dynorphin analogs such as those described in International Patent Application WO 96/06626 of Lee et al. and des-Tyr or des-Tyr-Gly analogs such as those disclosed in International Patent Application WO 93/25217 also of Lee et al. The dynorphis are highly potent opioids, and demonstrate selective affinity for the kappa receptor. See Goldstein, A., Peptides, Structure and Function, Proceedings of the 8th

American Peptides Symposium, Hruby, V.J. and Rich, D.H., eds., 409 (1983).

<u>Endorphins</u> – The endorphis are derived from the precursor protein -lipotropin. They have been found to elicit several biological reactions such as analgesia, behavioral changes and growth hormone release. See Akil, H. et al., Ann. Rev. Neurosci., 7, 223 (1984).

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Enkepalins & related peptides — Enkephalins and endorphins are neurohormones that inhibit transmission of pain impulses. The activity of neurons in both the central and peripheral nervous systems is affected by a large number of neurohormones that act on cells quite distant from their site of release. Neurohormones can modify the ability of nerve cells to respond to synaptic neurotransmitters. Several small peptides with profound effects on the nervous system have been discovered recently, for example enkephalins (e.g. Met-enkephalin and Leu-enkephalin) and endorphins (e.g. β endorphin). These three contain a common tetrapeptide sequence (Tyr-Gly-Gly-Phe) that is essential to their functions. Enkephalins and endorphins function as natural pain killers or opiates and decrease the pain responses in the central nervous system. See also Akil, H. etl al., Ann. Rev. Neurosci., 7, 223 (1984).

L. Thymic peptides (SEQ ID NOS: 928-934) – The thymus is thought to be responsible for the development and regulation of T cell immunity in both infants and adults. The thymus seems to exert its regulatory functions through the secretion of various noncellular, hormonelike products via its epithelial cells, called thymic peptides.

Thymic peptides are reported to have many effects on T cells. Several studies have reported that thymic peptides can assist development of immature, precursor cells into fully competent T cells. Thymic peptides seem to regulate the expression of various cytokine and monokine receptors on T cells and induce secretion of IL-2, interferon alpha, and interferon gamma (disease-fighting substances)

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when the immune system is challenged. There are reports that the use of thymic hormones in children with immuno-deficiencies caused by chemotherapy has resulted in an increase in circulating T cells, normalization of T cell subsets, and restoration of delayed hypersensitivity reactions.

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<u>Thymopoietin</u> - Thymopoietin is the largest of the known thymic hormones and consists of 49 amino acids.

<u>Thymulin</u> - Previously known as thymic serum factor, thymulin is the smallest of the chemically characterized thymic hormones and consists of 9 amino acids. Thymulin is the hormone responsible for stimulating the production of immune-system T cells

Thymopentin - Thymopentin is a small, synthesized thymic peptide drug, also known as therapeutic peptide-5 or Timunox. In the U.S. it is being developed as an AIDS therapy by the Immunobiology Research Institute. Thymopentin has been studied more extensively than most other thymic peptide drugs. At least one study has claimed a significant rise in T cells and slight clinical improvement in those patients who received thymopentin three times a week, compared to untreated control participants. Compared to the 14 untreated control participants, those taking the drug showed greater "immunologic stability" and some clinical improvement.

<u>Thymosin</u> - Thymosin is a mixture of 15 or more proteins. One of these proteins is thymosin alpha-1 which consists of 28 amino acids. Thymosin has therapeutic use for the treatment of primary immunodeficiencies and as a booster for influenza vaccine in renal dialysis patients. It is also being tested in ongoing clinical trials for activity against chronic hepatitis B and C, HIV infection, and certain forms of cancer.

Thymic Humoral Factor (THF) - THF is a thymic peptide currently being examined as an anti- HIV treatment. In preclinical

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studies in rats with CMV-related immunosuppression, THF restored immune competence through modulation of T cells. In addition, it may have therapeutic use in the treatment of herpes, causing (at least in one study) the viral infection's rapid regression and increase of T-cell populations.

L. Other Peptides

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Adrenomedullin Peptides (AM) (SEQ ID NOS: 935-945) -

Adrenomedullin is a potent vasodilator peptide that exerts major effects on cardiovascular functions. Its systemic administration causes a rapid and profound fall in blood pressure and an increase in pulmonary blood flow. Its other actions are bronchodilatation, being an inhibitor of drinking behavior and an inhibitor of angiotensin-induced aldosterone secretion.

See The Journal of Biological Chemistry, Vol. 270, No. 43, pp 25344-25347, 1995 and in the references cited therein

Allatostatin Peptides (SEQ ID NOS: 946-949) - Allatostatins are 6-18 amino acid peptides synthesized by insects to control production of juvenile hormones, which in turn regulate functions including metamorphosis and egg production. While neuropeptides of the allatostatin family inhibit *in vitro* production of juvenile hormone, which modulates aspects of development and reproduction in the cockroach, *Diploptera punctata*, they are susceptible to inactivation by peptidases in the hemolymph, gut, and bound to internal tissues.

Amyloid Beta-Protein Fragments (Aß fragments) (SEQ ID

NOS: 950-1010) – These are the principle component of the amyloid plaques that accumulate intracellularly and extracellularly in the neuritic plaques in the brain in Alzheimer's Disease. Aß is a 4.5 kD protein, about 40-42 amino acids long, that is derived from the C-terminus of amyloid precursor protein (APP). APP is a membrane-spanning glycoprotein that, in the normal processing pathway, is cleaved inside the Aß protein to produce α-sAPP, a secreted form of APP. Formation of α-sAPP precludes formation of Aß. It has been proposed that Aß

accumulates by virtue of abnormal processing of APP, so that compounds that inhibit the activity of enzymes responsible for Aß production are being sought. See, e.g., Wagner et al. Biotech. Report (1994/1995), pp. 106-107; and Selkoe (1993) TINS 16:403-409. Under certain conditions Aß peptides first aggregate and then are deposited as a folded ß-sheet structure that is characteristic of amyloid fibrils. ß-amyloid (1-42) forms aggregates at a significantly greater rate and to a greater extent than ß-amyloid (1-40).

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Antimicrobial peptides (SEQ ID NOS: 1011-1047) -

Antimicrobial peptides are a key component of the innate immune systems of most multicellular organisms, being active against one or more microorganisms such as bacteria, fungi, protozoa, yeast, and mycobacteria. Examples of such peptides include defensin, cecropin, buforin, and magainin. Despite broad divergences in sequence and taxonomy, most antimicrobial peptides share a common mechanism of action, *i.e.*. membrane permeabilization of the pathogen. They are classified in two broad groups: linear and cyclic. In the linear antimicrobial peptides, there are two subgroups: linear peptides tending to adopt α -helical amphipathic conformation and linear peptides of unusual composition, rich in amino acids such as Pro, Arg, or Trp. The cyclic group encompasses all cysteine-containing peptides, and can be further divided into two subgroups corresponding to single or multiple disulfide structures.

Most antimicrobial peptides provoke an increase in plasma membrane permeability. There is also evidence of other mechanisms, such as inhibition of specific membrane proteins, synthesis of stress proteins, arrest of DNA synthesis, breakage of single-strand DNA by defensins, interaction with DNA (without arrest of synthesis) by buforins, or production of hydrogen peroxide. Antimicrobial peptides can also act by triggering self-destructive mechanisms such as apoptosis in eukaryotic cells or autolysis in bacterial targets. Antimicrobial peptides are also known to act as inhibitors of enzymes produced by pathogenic

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organisms, either by serving as pseudo-substrates or by tight binding to the active sight that disturbs the access of the substrate.

Increased levels of antimicrobial peptides have been reported for several animal and human infections for example for α -defensins in *Mycobacterium, Pasteurella*, or *Cryptoporidium* infections and for a variety of peptides in blisters and wound fluid. Inflammatory situations or stimuli are also associated with induction of antibiotic peptides.

Depleted levels of antimicrobial peptides are associated to several pathologies. Thus, patients of specific granule-deficiency syndrome, completely lacking in α -defensins, suffer from frequent and severe bacterial infections. Low levels of histatins from saliva in HIV patients has been correlated with a higher incidence of oral candidiasis and fungal infections. Perhaps the most compelling illustration of the implication of antimicrobial peptides in human pathology comes from cystic fibrosis, a genetic disease associated with recurrent bacterial infections of the airways. The defective chloride channel causing the disease increases the salinity of the alveolar fluid, and thus impairs the bactericidal activity of β -defensins, which are salt sensitive. Andreu D, (Ed.)(1998) "Antimicrobial peptides" Biopolymers (Peptide Science) vol 47, N° 6, pp413-491. A. Andreu, L. Rivas (1998) Animal Antimicrobial Peptides: An Overview, Biopolymers (Pep. Sci.) 47: pp415-433.

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Antioxidant Peptides (SEQ ID NOS: 1048-1050) - Antioxidants are agents that prevents oxidative damage to tissue. Mammalian cells are continuously exposed to activated oxygen species such as superoxide, hydrogen peroxide, hydroxyl radical, and singlet oxygen. These reactive oxygen intermediates are generated *in vivo* by cells in response to aerobic metabolism, catabolism of drugs and other xenobiotics, ultraviolet and x-ray radiation, and the respiratory burst of phagocytic cells (such as white blood cells) to kill invading bacteria such as those introduced through wounds. Hydrogen peroxide, for example,

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is produced during respiration of most living organisms especially by stressed and injured cells.

One example of antioxidant peptides is natural killer-enhancing factor B (NKEF-B), which belongs to a highly conserved family of recently discovered antioxidants. Natural killer-enhancing factor (NKEF) was identified and cloned on the basis of its ability to increase NK cytotoxicity. Two genes, NKEF-A and -B, encode NKEF proteins and sequence analysis presented suggests that each belongs to a highly conserved family of antioxidants. The role of NKEF-B as an antioxidant has been demonstrated by its protection of transfected cells to oxidative damage by hydrogen peroxide. NKEF-B has antioxidant activities toward prooxidants such as alkyl hydroperoxide and MeHg. Together with its antioxidant activity, the induction of NKEF-B by HP indicates that NKEF-B is an important oxidative stress protein providing protection against a variety of xenobiotic toxic agents.

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Apoptosis Related Peptides (SEQ ID NOS: 1051-1075) -

Animal cells can self-destruct via an intrinsic program of cell death (Steller, 1995). Apoptosis is a form of programmed cell death that is characterized by specific morphologic and biochemical properties (Wyllie et al., 1980). Morphologically, apoptosis is characterized by a series of structural changes in dying cells: blebbing (i.e. blistering) of the plasma membrane, condensation of the cytoplasm and nucleus, and cellular fragmentation into membrane apoptotic bodies (Steller, 1995; Wyllie et al., 1980).

Biochemically, apoptosis is characterized by the degradation of chromatin, initially into large fragments of 50-300 kilobases and subsequently into smaller fragments that are monomers and multimers of 200 bases (Oberhammer *et al.*, 1993; Wyllie, 1980). Other biochemical indicators of apoptosis are induced or increased levels of the protein clusterin (Pearse *et al.*, 1992), also known as TRPM-2 or SGP-2, and activation of the enzyme typell transglutaminase, which

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crosslinks proteins to the envelope of apoptotic bodies (Fesus *et al.*, 1991). Apoptosis is a complex phenomenon of related morphological and biochemical processes that can vary with tissue and cell type (Zakeri *et al.*, 1995).

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The execution of apoptosis minimizes the leakage of cellular constituents from dying cells (apoptosis causes the cell to involute). For example, proteases could damage adjacent cells or stimulate an inflammatory response. This cardinal feature of apoptosis distinguishes it from necrosis, which usually results from trauma that causes injured cells to swell and lyse, releasing the cytoplasmic material that stimulates an inflammatory response (Steller, 1995; Wyllie *et al.*, 1980)

Bag Cell Peptides (BCP) (SEQ ID NOS: 1076-1080) – The neuropeptidergic bag cells of the marine mollusc *Aplysia californica* are involved in the egg-laying behavior of the animal. These neurosecretory cells synthesize an egg-laying hormone (ELH) precursor protein, yielding multiple bioactive peptides, including ELH, several bag cell peptides (BCP) and acidic peptide (AP). The bag cells of the marine mollusc *Aplysia californica* are well-characterized neuroendocrine cells that initiate egg laying. During sexual maturation, these cells (bag cell neurons), develop the capability of storing hormones that are released during periods of nervous system stimulation. The hormones are important to the process of egg laying, and so must not be released before the animal is sexually mature. Alpha-bag cell peptide belong to a small family of structurally related peptides that can elicit bag-cell activity in vitro.

Bombesin (SEQ ID NOS: 1081-1090) – Bombesin is a bioactive tetradecapeptide neuropeptide that belongs to a family of peptides sharing a common C terminal sequence, Trp-Ala-X-Gly-His-Met-NH2, and the N terminal region. It has a modulatory role found in nerves of the brain and gut that prevents gastric injury by release of endogenous gastrin. The mammalian homologue of bombesin is gastrin-releasing peptide (GRP).

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Bone Gla Protein Peptides (SEQ ID NOS: 1091-1097) -

Osteocalcin (bone Gla-protein, or BGP) is produced and secreted by osteoblasts in the process of bone formation. As with collagen, this protein is a component of bone matrix. Serum osteocalcin rises when bone formation rates increase. Levels are high during puberty when bone growth is most rapid. Often levels are also high in diseases having high bone turnover, such as hyperparathyroidism and hyperthyroidism. In postmenopausal osteoporosis, osteocalcin levels are sometimes increased, reflecting the increased turnover of bone secondary to rapid bone resorption. In senile osteoporosis, occurring in more elderly subjects, osteocalcin levels are more likely to be low, reflecting reduced rates of both bone turnover and bone formation. A treatment regimen that increases bone formation also raises the serum osteocalcin levels.

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CART Peptides (SEQ ID NOS: 1098-1100) - Cocaine and amphetamine regulated transcript peptide (CART), is a recently discovered hypothalamic peptide with a potent appetite suppressing activity. In the rat the CART gene encodes a peptide of either 129 or 116 amino acid residues whereas only the short form exists in humans. The predicted signal sequence is 27 amino acid residues resulting in a prohormone of 102 or 89 residues. The C-terminal end of CART, consisting of 48 amino acid residues and 3 disulphide bonds, is thought to constitute a biologically active part of the molecule.

In the central nervous system CART is highly expressed in many hypothalamic nuclei, some of which are involved in regulating feeding behavior. The CART mRNA is regulated by leptin, and the expressed CART is a potent inhibitor of feeding that even overrides the feeding response induced by neuropeptide Y. The putative CART receptor is therefore a potential therapeutic target for an anti-obesity drug. See CART, a new anorectic peptide Thim L; Kristensen P; Larsen PJ; Wulff BS, Int J Biochem Cell Biol, 30(12):1281-4 1998 Dec.

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Cell Adhesion Peptides (SEQ ID NO: 1101) - Cellular adhesion peptides are directly involved in the cellular response to external stimuli. For example, during an inflammatory response, leukocytes must leave the plasma compartment and migrate to the point of antigenic insult. The mechanism of this migratory event is a complex interplay between soluble mediators and membrane-bound cellular adhesion molecules. Soluble cellular chemotactic factors, which are produced in the damaged tissue by a variety of resident cells, set up a chemical concentration gradient out to the plasma compartment. Interaction of these factors with their receptors on leukocytes leads to a directional migration of the leukocytes toward increasing concentrations of the chemotactic factor. Simultaneously, various adhesion peptides are upregulated on the leukocyte which mediate the initial rolling on the endothelial tissue, binding to a specific ligand on the activated endothelial tissue, and finally migration between endothelial cells into the tissue. The steps in this cascade of events are mediated by the interaction of specific cell surface proteins, termed "cell adhesion molecules such as. E-selectin (ELAM-1, endothelial leukocyte adhesion molecule-1), ICAM-1 (intercellular adhesion molecule-1), and VCAM-1 (vascular cell adhesion molecule-1).

Chemotactic Peptides (SEQ ID NOS: 1102-1113) — Chemotactic peptides are peptides that stimulate the migration of white cells, leukocytes and macrophages into tissues at the site of infection or injury or alternatively the prevent the migration of these same cells away from these sites.

Complement Inhibitors (SEQ ID NOS: 1114-1120) - Inhibition of complement attack on xenotransplants may be accomplished by the use of complement inhibitors. The rejection of transplanted organs may involve both an extremely rapid hyperacute rejection (HAR) phase and a slower cellular rejection phase. HAR of xenotransplants is initiated by preformed "natural" antibodies that bind to donor organ endothelium and activate complement attack by the recipient immune system. Activation of complement leads to the generation of fluid phase (C3a, C5a) and

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membrane bound (C3b and C5b-9, i.e., C5b, C6, C7, C8, and C9) proteins with chemotactic, procoagulant, proinflammatory, adhesive, and cytolytic properties. Complement inhibitors inhibit this process.

Cortistatin Peptides (SEQ ID NOS: 1121-1124) - Cortistatin, whose mRNA accumulates during sleep deprivation, apparently acts by 5 antagonizing the effects of acetylcholine on cortical excitability, thereby causing synchronization brain slow waves. Cortistatin-14 (CST-14) shares 11 of its 14 residues with somatostatin-14 (SRIF-14), yet its effects on sleep physiology, locomotor behavior and hippocampal function are quite different from those of somatostatin.

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Fibronectin Fragments & Fibrin Related Peptides (SEQ ID NOS: 1125-1174) - Fibronectin is a large glycoprotein that is composed of blocks of three types of repeating, homologous peptide sequences. Several of the homologous blocks form functional domains that are organized in a linear array on two nearly identical subunit arms. Each arm can be divided into functional domains, which are often referred to by one of the substances which bind in that region, for example the heparin-binding fragment, the fibrin binding fragment, and the cellbinding fragment. In several cell types, the Arg-Gly-Asp (RGD) sequence in the cell-binding domain of fibronectin interacts with a cellsurface glycoprotein designated lib/IIIa. Fibronectin also binds to extracellular and basement-membrane components, to the envelope glycoprotein of viruses, to a variety of bacteria including staphylococci and streptococci, and to parasites such as Trypanosoma cruzi and Leishmania species.

Fibronectin has several adhesive functions, for example cell-tocell adhesion, cell-to-basement-membrane attachment, and clot stabilization. In addition, fibronectin promotes embryogenesis, nerve regeneration, fibroblast migration, macrophage function, and pathogen (virus, fungus, bacteria, and protozoa) binding to mammalian cells and extracellular matrix. Thus, fibronectin is involved in the pathogenesis of 5

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infections from the initiation of the infection through the final stages of wound healing. <u>See</u> Proctor, R.A., Rev. Infect. Dis., 9, 317 (1987).

FMRF and Analog peptides (SEQ ID NOS: 1175-1187) –

FMRF are neuropeptides encoded in the FMRFamide gene and have a common C-terminal FMRFamide but different N-terminal extensions.

FMRFamide-related peptides (FaRPs) are present throughout the animal kingdom and affect both neural and gastrointestinal functions.

Organisms have several genes encoding numerous FaRPs with a common C-terminal structure but different N-terminal amino acid extensions.

Galanin & related peptides (SEQ ID NOS: 1188-1208)- Galanin is a 29-30 amino acid peptide originally isolated from pig small intestine. It is found in two biologically active forms: GAL (1-19), and GAL (1-30), a non-amidated form. It has many biological roles including: the inhibition of the release of biogenic amines in the hypothalamus, the pre- and post-synaptic inhibition of cholinergic function, the maintenance of gastrointestinal homeostasis, and the regulation of insulin and glucagon secretion.

Growth Factors & related peptides (SEQ ID NOS: 1209-1240)

– Growth factors are a family of proteins that regulate cell division.
Some growth factors are cell type specific, stimulating division of only those cells with appropriate receptors. Other growth factors are more general in their effects. There are also extracellular factors that antagonize the effects of growth factors, slowing or preventing division (for example transforming growth factor beta and tumor necrosis factor). These extracellular signals act through cell surface receptors very similar to those for hormones, and by similar mechanisms: the production of intracellular second messangers, protein phosphorylation, and ultimately, alteration of gene expression.

Gtherapeutic peptide-Binding protein fragments (SEQ ID NOS: 1241-1246) – Members of a family of Gtherapeutic peptide-binding regulatory proteins (G-proteins) transduce signals from

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membrane-bound receptors to intracellular effectors. The family includes G_s and G_i, which are responsible for stimulation and inhibition, respectively, of adenylate cyclase. Transducin (T), localized in the disc membranes of retinal rod outer segments, couples activation of rhodopsin by light to increased cyclinc GMP phosphodiesterase activity.

**Go, found originally in bovine brain, is a fourth member of the family.

Purified G proteins have similar physical properties. They are heterodimers composed of α , β , and γ subunits. The α subunits bind and hydrolyze Gtherapeutic peptide. See S. M. Mumby et al., PNAS 83, 265 (1986) and Lehninger p. 764.

Guanylin and Uroguanylin (SEQ ID NOS: 1247-1249) -

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Guanylin and uroguanylin are peptides isolated from intestinal mucosa, and urine, which regulate cyclic GMP production in enterocytes bind to and activate guanylate cyclase C and control salt and water transport in many epithelia in vertebrates, mimicking the action of several heat-stable bacteria enterotoxins. In the kidney, both of them have well-documented natriuretic and kaliuretic effects.

Chloride secretion in the intestine is regulated by these hormones via activation of guanylate cyclase C (GC-C). Both peptides are expressed in a variety of tissues and organs, including the kidney. In the isolated perfused kidney and in vivo these hormones induce natriuresis and diuresis, however, localisation and cellular mechanisms of their action in the kidney are still unknown.

Inhibin Peptides (SEQ ID NOS: 1250-1255) – Inhibin is composed of two subunits (α is 134 amino acids; β is 115 and 116 amino acids). Its role is inhibition of FSH secretion. The two inhibin isoforms, inhibin A and inhibin B, are produced by the gonads in the course of gamete maturation and have different patterns of secretion during the menstrual cycle. Inhibins are also produced by the placenta and fetal membranes and may be involved in physiological adaptation of pregnancy. Clinically, inhibins may serve as sensitive tumor markers in postmenopausal women, or as useful tools for evaluating ovarian

reserve in infertile women; they may also be used in the diagnosis of materno-fetal disorders and to prevent maturation of the ovum or to inhibit ovulation.

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Interleukin (IL) and Interleukin Receptor Proteins (SEQ ID NOS: 1256-1263) – Interleukins are growth factors targeted to cells of hematopoietic origin. A variety of biological activities associated with immune and inflammatory responses have been ascribed to interleukins. These responses include fever, cartilage breakdown, bone resorption, thymocyte proliferation, activation of T and B lymphocytes, induction of acute-phase protein synthesis from hepatocytes, fibroblast proliferation, and differentiation and proliferation of bone marrow cells.

Laminin Fragments (SEQ ID NOS: 1264-1284) - Laminin, the major noncollagenous glycoprotein of basement membranes, has been shown to promote the adhesion, spreading, and migration of a variety of tumor cell types in vitro. In particular, the major current studies in the laboratory utilize intact laminin, purified proteolytic fragments of laminin, and synthetic peptides of laminin to identify functionally active sites on this large protein. Components of such basement membranes are important modulators of growth, development, and differentiation for various cell types. A conjugated laminin could be used to prevent inflamation or fibrosis in tissues.

This category also includes the peptide kringle-5 (or K-5). As used herein, the term "kringle 5" refers to the region of mammalian plasminogen having three disulfide bonds which contribute to the specific three-dimensional confirmation defined by the fifth kringle region of the mammalian plasminogen molecule. One such disulfide bond links the cysteine residues located at amino acid positions 462 and 541, a second links the cysteine residues located at amino acid positions 483 and 524 and a third links the cysteine residues located at amino acid positions 512 and 536. The term "kringle 5 peptide peptides" refers to peptides with anti-angiogenic activity of between 4 and 104 amino acids

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(inclusive) with a substantial sequence homology to the corresponding peptide fragment of mammalian plasminogen.

Leptin Fragment Peptides (SEQ ID NOS: 1285-1288) – Leptin, the protein product of the obesity gene, is secreted by fat cells. Leptin is involved in the regulation of bodyweight and metabolism in man and might also be involved in the pathophysiology of the insulin resistance syndrome, which is associated with the development of cardiovascular diseases

<u>Leucokinins (SEQ ID NOS: 1289-98)</u> - Leucokinins are a group of widespread insect hormones that stimulate gut motility and tubule fluid secretion rates. In tubules, their major action is to raise chloride permeability by binding to a receptor on the basolateral membrane.

Pituitary Adenylate Cyclase Activating Polypeptide (PACAP) (SEQ ID NOS: 1299-1311) — It is a thirty-eight amino acid peptide first isolated from ovine hypothalamus, which also occurs in a 27 amino acid form called PACAP-27. PACAP has been localized in the hypothalamus, elsewhere in the brain, respiratory tract and gastrointestinal system. It has many biological actions, including neurotransmitter and hormonal functions, involvement in regulation of energy metabolism, and neuronal cytoprotective activity.

Pancreastatin (SEQ ID NOS: 1312-1324) — Pancreastatin is a 49 amino acid peptide first isolated, purified and characterized from porcine pancreas. Its biological activity in different tissues can be assigned to the C-terminal part of the molecule. Pancreastatin has a prohormonal precursor, chromogranin A, which is a glycoprotein present in neuroendocrine cells, including the endocrine pancreas

<u>Polypeptides (SEQ ID NOS: 1325-1326)</u> – these are repetitive chains. Two examples are provided: (pro-Hyp-Gly)10*20H20 and Poly-L-Lysine Hydrochloride.

30 <u>Signal Transduction Reagents (SEQ ID NOS: 1327-1367)</u> – Signal transduction is the process by which an extracellular signal (for example chemical, mechanical, or electrical) is amplified and converted

to a cellular response. Many reagents are involved in this process, for example achatin-1, glycogen synthase, autocamtide 2, calcineurin autoinhibitory peptide, calmodulin dependent protein kinase II, calmodulin dependent protein kinase substrate, calmodulin dependent protein kinase substrate analog, CKS-17, Cys-Kemptide, autocamtide 2, malantide, melittin, phosphate acceptor peptide, protein kinase C fragments, P34cd2 kinase fragment, P60c-src substrate II, protein kinase A fragments, tyrosine protein kinase substrate, syntide 2, S6 kinase substrate peptide 32, tyrosine specific protein kinase inhibitor, and their derivatives and fragments.

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Thrombin Inhibitors (SEQ ID NOS: 1368-1377) - Thrombin is a key regulatory enzyme in the coagulation cascade; it serves a pluralistic role as both a positive and negative feedback regulator. In addition to its direct effect on hemostasis, thrombin exerts direct effects on diverse cell types that support and amplify pathogenesis of arterial thrombus disease. The enzyme is the strongest activator of platelets causing them to aggregate and release substances (eg. ADP TXA.sub.2 NE) that further propagate the thrombotic cycle. Platelets in a fibrin mesh comprise the principal framework of a white thrombus. Thrombin also exerts direct effects on endothelial cells causing release of vasoconstrictor substances and translocation of adhesion molecules that become sites for attachment of immune cells. In addition, the enzyme causes mitogenesis of smooth muscle cells and proliferation of fibroblasts. From this analysis, it is apparent that inhibition of thrombin activity by thrombin inhibitors constitutes a viable therapeutic approach towards the attenuation of proliferative events associated with thrombosis.

<u>Toxins (SEQ ID NOS: 1378-1415)</u> — A toxin can be conjugated using the present invention to target cancer cells, receptors, viruses, or blood cells. Once the toxin binds to the target cells the toxin is allowed to internalize and cause cell toxicity and eventually cell death. Toxins have been widely used as cancer therapeutics.

One example of a class of toxins is the mast cell degranulating peptide, a cationic 22-amino acid residue peptide with two disulfide bridges isolated from bee venom, causes mast cell degranulation and histamine release at low concentrations and has anti-inflammatory activity at higher concentrations. It is a powerful anti-inflammatory, more than 100 times more effective than hydrocortisone in reducing inflammation. Because of these unique immunologic properties, MCD peptide may serve as a useful tool for studying secretory mechanisms of inflammatory cells such as mast cells, basophils, and leukocytes, leading to the design of compounds with therapeutic potential. An example of a mast cell degranulating peptide is mastoparans, originating from wasp venom. It degranulates mast cells in the concentration of 0.5 µg/ml and releases histamine from the cells in the same concentration. See IY. Hirai et al., Chem. Pharm. Bull. 27, 1942 (1979).

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Other examples of such toxins include omega-agatoxin TK, agelenin, apamin, calcicudine, calciseptine, charbdotoxin, chlorotoxin, conotoxins, endotoxin inhibitors, gegraphutoxins, iberiotoxin, kaliotoxin, mast cell degranulating peptides, margatoxin, neurotoxin NSTX-3, PLTX-II, scyllatoxin, stichodactyla toxin, and derivatives and fragments thereof.

<u>Trypsin Inhibitors (SEQ ID NOS: 1416-1418)</u> - Trypsin inhibitors functions as an inhibitors of trypsin, as well as other serine proteases. Useful for treatment of lung inflammation, pancreatitis, myocardial infarction, cerebrovascular ischemia

Virus Related Peptides (SEQ ID NOS: 1419-1529) — Virus related peptides are proteins related to viruses, for example virus receptors, virus inhibitors, and envelope proteins. Examples include but are not limited to peptide inhibitors of human immunodeficiency virus (HIV), respiratory syncytial virus (RSV), human parainfluenza virus (HPV), measles virus (MeV), and simian immunodeficiency virus (SIV),

fluorogenic Human CMV Protease Substrate, HCV Core Protein, HCV NS4A Protein, Hepatitis B Virus Receptor Binding Fragment, Hepatitis B Virus Pre-S Region, Herpes Virus Inhibitor 2, HIV Envelope Protein Fragment, HIV gag fragment, HIV substrate, HIV-1 Inhibitory Peptide, peptide T, T21, V3 decapeptide, Virus Replication Inhibitor Peptide, and their fragments and derivatives.

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These peptides can be administered therapeutically. For example, peptide T is a chain of 8 amino acids from the V2 region of HIV-1 gp120. These amino acids look like a portion of HIV's outer envelope. It is under investigation as a treatment for HIV-related neurological and constitutional symptoms, as peptide T may be able to alleviate symptoms like fevers, night sweats, weight loss, and fatigue. It has also been shown to resolve psoriatic lesions.

Miscellaneous peptides (SEQ ID NOS: 1529-1617) - Including adjuvant peptide analogs, alpha mating factor, antiarrhythmic peptide, anorexigenic peptide, alpha-1 antitrypsin, bovine pineal antireproductive peptide, bursin, C3 peptide P16, cadherin peptide, chromogranin A fragment, contraceptive tetrapeptide, conantokin G, conantokin T, crustacean cardioactive peptide, C-telopeptide, cytochrome b588 peptide, decorsin, delicious peptide, delta-sleep-inducing peptide, diazempam-binding inhibitor fragment, nitric oxide synthase blocking peptide, OVA peptide, platelet calpain inhibitor (P1), plasminogen activator inhibitor 1, rigin, schizophrenia related peptide, sodium potassium Atherapeutic peptidease inhibitor-1, speract, sperm activating peptide, systemin, thrombin receptor agonist (three peptides), tuftsin, adipokinetic hormone, uremic pentapeptide, Antifreeze Polypeptide, tumor necrosis factor, leech [Des Asp10]Decorsin, L-Ornithyltaurine Hydrochloride, p-Aminophenylacetyl Tuftsin, Ac-Glu-Glu-Val-Val-Ala-Cys-pNA, Ac-Ser-Asp-Lys-Pro, Ac-rfwink-NH2, Cys-Gly-Tyr-Gly-Pro-Lys-Lys-Lys-Arg-Lys-Val-Gly-Gly, DAla-Leu, D-D-D-D, D-D-D-D-D, N-P-N-A-N-P-N-A, V-A-I-T-V-L-V-K, V-G-V-R-V-R, V-I-H-S, V-P-D-P-R, Val-Thr-Cys-Gly, R-S-R, Sea Urchin Sperm Activating Peptide, SHU-

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9119 MC3-R & MC4-R Antagonist, glaspimod (immunostimulant, useful against bacterial infections, fungal infections, immune deficiency immune disorder, leukopenia), HP-228 (melanocortin, useful against chemotherapy induced emesis, toxicity, pain, diabetes mellitus, inflammation, rheumatoid arthritis, obesity), alpha 2-plasmin inhibitor (plasmin inhibitor), APC tumor suppressor (tumor suppressor, useful against neoplasm), early pregnancy factor (immunosuppressor), endozepine diazepam binding inhibitor (receptor peptide), gamma interferon (useful against leukemia), glandular kallikrei n-1 (immunostimulant), placental ribonuclease inhibitor, sarcolecin binding protein, surfactant protein D, wilms' tumor suppressor, wilm's tumor suppressor, GABAB 1b receptor peptide, prion related peptide (iPrP13), choline binding protein fragment (bacterial related peptide), telomerase inhibitor, cardiostatin peptide, endostatin derived peptide (angiogenesis inhibitor), prion inhibiting peptide, N-methyl D-aspartate receptor antagonist, C-peptide analog (useful against diabetic complications).

2. Modified Therapeutic Peptides

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This invention relates to modified therapeutic peptides and their derivatives. The modified therapeutic peptides of the invention include reactive groups which can react with available reactive functionalities on blood components to form covalent bonds. The invention also relates to such modifications, such combinations with blood components and methods for their use. These methods include extending the effective therapeutic *in vivo* half life of the modified therapeutic peptides.

To form covalent bonds with functionalities on a protein, one may use as a reactive group a wide variety of active carboxyl groups, particularly esters, where the hydroxyl moiety is physiologically acceptable at the levels required to modify the therapeutic peptide. While a number of different hydroxyl groups may be employed in these linking agents, the most convenient would be N-hydroxysuccinimide (NHS), and N-hydroxy-sulfosuccinimide (sulfo-NHS).

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Primary amines are the principal targets for NHS esters as diagramed in schematic 1A below. Accessible α -amine groups present on the N-termini of proteins react with NHS esters. However, α -amino groups on a protein may not be desirable or available for the NHS coupling. While five amino acids have nitrogen in their side chains, only the ϵ -amine of lysine reacts significantly with NHS esters. An amide bond is formed when the NHS ester conjugation reaction reacts with primary amines releasing N-hydroxysuccinimide as demonstrated in schematic 1A below.

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Schematic 1A

In the preferred embodiments of this invention, the functionality on the protein will be a thiol group and the reactive group will be a maleimido-containing group such as gamma-maleimide-butyralamide (GMBA) or MPA. The maleimido group is most selective for sulfhydryl groups on peptides when the pH of the reaction mixture is kept between 6.5 and 7.4 as shown in schematic 1B below. At pH 7.0, the rate of reaction of maleimido groups with sulfhydryls is 1000-fold faster than with amines. A stable thioether linkage between the maleimido group and the sulfhydryl is formed which cannot be cleaved under physiological conditions.

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The therapeutic peptides and peptide derivatives of the invention may be modified for specific labeling and non-specific labeling of blood components.

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A. Specific Labeling

Preferably, the therapeutic peptides of this invention are designed to specifically react with thiol groups on mobile blood proteins. Such reaction is preferably established by covalent bonding of a therapeutic peptide modified with a maleimide link (e.g. prepared from GMBS, MPA or other maleimides) to a thiol group on a mobile blood protein such as serum albumin or IgG.

Under certain circumstances, specific labeling with maleimides offers several advantages over non-specific labeling of mobile proteins with groups such as NHS and sulfo-NHS. Thiol groups are less abundant *in vivo* than amino groups. Therefore, the maleimide derivatives of this invention will covalently bond to fewer proteins. For example, in albumin (the most abundant blood protein) there is only a single thiol group. Thus, therapeutic peptide-maleimide-albumin conjugates will tend to comprise approximately a 1:1 molar ratio of therapeutic peptide to albumin. In addition to albumin, IgG molecules (class II) also have free thiols. Since IgG molecules and serum albumin make up the majority of the soluble protein in blood they also make up

the majority of the free thiol groups in blood that are available to covalently bond to maleimide-modified therapeutic peptides.

Further, even among free thiol-containing blood proteins, specific labeling with maleimides leads to the preferential formation of therapeutic peptide-maleimide-albumin conjugates, due to the unique characteristics of albumin itself. The single free thiol group of albumin, highly conserved among species, is located at amino acid residue 34 (Cys³⁴). It has been demonstrated recently that the Cys³⁴ of albumin has increased reactivity relative to free thiols on other free thiolcontaining proteins. This is due in part to the very low pK value of 5.5 for the Cvs³⁴ of albumin. This is much lower than typical pK values for cysteines residues in general, which are typically about 8. Due to this low pK, under normal physiological conditions Cys³⁴ of albumin is predominantly in the ionized form, which dramatically increases its reactivity. In addition to the low pK value of Cys³⁴, another factor which enhances the reactivity of Cvs³⁴ is its location, which is in a crevice close to the surface of one loop of region V of albumin. This location makes Cys³⁴ very available to ligands of all kinds, and is an important factor in Cys³⁴'s biological role as free radical trap and free thiol scavenger. These properties make Cys³⁴ highly reactive with therapeutic peptidemaleimides, and the reaction rate acceleration can be as much as 1000fold relative to rates of reaction of therapeutic peptide-maleimides with other free-thiol containing proteins.

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Another advantage of therapeutic peptide-maleimide-albumin conjugates is the reproducibility associated with the 1:1 loading of peptide to albumin specifically at Cys³⁴. Other techniques, such as glutaraldehyde, DCC, EDC and other chemical activations of, for example, free amines lack this selectivity. For example, albumin contains 52 lysine residues, 25-30 of which are located on the surface of albumin and accessible for conjugation. Activating these lysine residues, or alternatively modifying peptides to couple through these lysine residues, results in a heterogenous population of conjugates.

Even if 1:1 molar ratios of peptide to albumin are employed, the yield will consist of multiple conjugation products, some containing 0, 1, 2 or more peptides per albumin, and each having peptides randomly coupled at any one of the 25-30 available lysine sites. Given the numerous combinations possible, characterization of the exact composition and nature of each batch becomes difficult, and batch-to-batch reproducibility is all but impossible, making such conjugates less desirable as a therapeutic. Additionally, while it would seem that conjugation through lysine residues of albumin would at least have the advantage of delivering more therapeutic agent per albumin molecule, studies have shown that a 1:1 ratio of therapeutic agent to albumin is preferred. In an article by Stehle, et al., "The Loading Rate Determines Tumor Targeting Properties of Methotrexate-Albumin Conjugates in Rats," Anti-Cancer Drugs, Vol. 8, pp. 677-685 (1997), incorporated herein in its entirety, the authors report that a 1:1 ratio of the anti-cancer methotrexate to albumin conjugated via glutaraldehyde gave the most promising results. These conjugates were taken up by tumor cells, whereas conjugates bearing 5:1 to 20:1 methotrexate molecules had altered HPLC profiles and were quickly taken up by the liver in vivo. It is postulated that at these higher ratios, conformational changes to albumin diminish its effectiveness as a therapeutic carrier.

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Through controlled administration of maleimide-therapeutic peptides *in vivo*, one can control the specific labeling of albumin and IgG *in vivo*. In typical administrations, 80-90% of the administered maleimide-therapeutic peptides will label albumin and less than 5% will label IgG. Trace labeling of free thiols such as glutathione will also occur. Such specific labeling is preferred for *in vivo* use as it permits an accurate calculation of the estimated half-life of the administered agent.

In addition to providing controlled specific *in vivo* labeling, maleimide-therapeutic peptides can provide specific labeling of serum albumin and IgG *ex vivo*. Such *ex vivo* labeling involves the addition of maleimide-therapeutic peptides to blood, serum or saline solution

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containing serum albumin and/or IgG. Once modified *ex vivo* with maleimide-therapeutic peptides, the blood, serum or saline solution can be readministered to the blood for *in vivo* treatment.

In contrast to NHS-peptides, maleimide-therapeutic peptides are generally quite stable in the presence of aqueous solutions and in the presence of free amines. Since maleimide-therapeutic peptides will only react with free thiols, protective groups are generally not necessary to prevent the maleimide-therapeutic peptides from reacting with itself. In addition, the increased stability of the peptide permits the use of further purification steps such as HPLC to prepare highly purified products suitable for *in vivo* use. Lastly, the increased chemical stability provides a product with a longer shelf life.

B. Non-Specific Labeling

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The therapeutic peptides of the invention may also be modified for non-specific labeling of blood components. Bonds to amino groups will generally be employed, particularly with the formation of amide bonds for non-specific labeling. To form such bonds, one may use as a chemically reactive group coupled to the therapeutic peptide a wide variety of active carboxyl groups, particularly esters, where the hydroxyl moiety is physiologically acceptable at the levels required. While a number of different hydroxyl groups may be employed in these linking agents, the most convenient would be N-hydroxysuccinimide (NHS) and N-hydroxy-sulfosuccinimide (sulfo-NHS).

Other linking agents which may be utilized are described in U.S. Patent 5,612,034, which is hereby incorporated herein.

The various sites with which the chemically reactive groups of the non-specific therapeutic peptides may react *in vivo* include cells, particularly red blood cells (erythrocytes) and platelets, and proteins, such as immunoglobulins, including IgG and IgM, serum albumin, ferritin, steroid binding proteins, transferrin, thyroxin binding protein, α -2-macroglobulin, and the like. Those receptors with which the derivatized

therapeutic peptides react, which are not long-lived, will generally be eliminated from the human host within about three days. The proteins indicated above (including the proteins of the cells) will remain in the bloodstream at least three days, and may remain five days or more (usually not exceeding 60 days, more usually not exceeding 30 days) particularly as to the half life, based on the concentration in the blood.

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For the most part, reaction will be with mobile components in the blood, particularly blood proteins and cells, more particularly blood proteins and erythrocytes. By "mobile" is intended that the component does not have a fixed situs for any extended period of time, generally not exceeding 5 minutes, more usually one minute, although some of the blood components may be relatively stationary for extended periods of time. Initially, there will be a relatively heterogeneous population of labeled proteins and cells. However, for the most part, the population within a few days after administration will vary substantially from the initial population, depending upon the half-life of the labeled proteins in the blood stream. Therefore, usually within about three days or more, lgG will become the predominant labeled protein in the blood stream.

Usually, by day 5 post-administration, IgG, serum albumin and erythrocytes will be at least about 60 mole %, usually at least about 75 mole %, of the conjugated components in blood, with IgG, IgM (to a substantially lesser extent) and serum albumin being at least about 50 mole %, usually at least about 75 mole %, more usually at least about 80 mole %, of the non-cellular conjugated components.

The desired conjugates of non-specific therapeutic peptides to blood components may be prepared *in vivo* by administration of the therapeutic peptides directly to the patient, which may be a human or other mammal. The administration may be done in the form of a bolus or introduced slowly over time by infusion using metered flow or the like.

If desired, the subject conjugates may also be prepared *ex vivo* by combining blood with modified therapeutic peptides of the present invention, allowing covalent bonding of the modified therapeutic peptides

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to reactive functionalities on blood components and then returning or administering the conjugated blood to the host. Moreover, the above may also be accomplished by first purifying an individual blood component or limited number of components, such as red blood cells, immunoglobulins, serum albumin, or the like, and combining the component or components *ex vivo* with the chemically reactive Itherapeutic peptides. The labeled blood or blood component may then be returned to the host to provide *in vivo* the subject therapeutically effective conjugates. The blood also may be treated to prevent coagulation during handling *ex vivo*.

3. Synthesis of Therapeutic Peptides Used in the Present Invention

Peptide fragments may be synthesized by standard methods of solid phase peptide chemistry known to those of ordinary skill in the art. For example, peptide fragments may be synthesized by solid phase chemistry techniques following the procedures described by Steward and Young (Steward, J. M. and Young, J. D., Solid Phase Peptide Synthesis, 2nd Ed., Pierce Chemical Company, Rockford, Ill., (1984) using an Applied Biosystem synthesizer. Similarly, multiple fragments may be synthesized then linked together to form larger fragments. These synthetic peptide fragments can also be made with amino acid substitutions at specific locations.

For solid phase peptide synthesis, a summary of the many techniques may be found in J. M. Stewart and J. D. Young, Solid Phase Peptide Synthesis, W. H. Freeman Co. (San Francisco), 1963 and J. Meienhofer, Hormonal Proteins and Peptides, vol. 2, p. 46, Academic Press (New York), 1973. For classical solution synthesis see G. Schroder and K. Lupke, The Peptides, Vol. 1, Acacemic Press (New York). In general, these methods comprise the sequential addition of one or more amino acids or suitably protected amino acids to a growing peptide chain. Normally, either the amino or carboxyl group of the first

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amino acid is protected by a suitable protecting group. The protected or derivatized amino acid is then either attached to an inert solid support or utilized in solution by adding the next amino acid in the sequence having the complimentary (amino or carboxyl) group suitably protected and under conditions suitable for forming the amide linkage. The protecting group is then removed from this newly added amino acid residue and the next amino acid (suitably protected) is added, and so forth.

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After all the desired amino acids have been linked in the proper sequence, any remaining protecting groups (and any solid support) are removed sequentially or concurrently to afford the final polypeptide. By simple modification of this general procedure, it is possible to add more than one amino acid at a time to a growing chain, for example, by coupling (under conditions which do not racemize chiral centers) a protected tripeptide with a properly protected dipeptide to form, after deprotection, a pentapeptide.

A particularly preferred method of preparing compounds of the present invention involves solid phase peptide synthesis wherein the amino acid α -N-terminal is protected by an acid or base sensitive group. Such protecting groups should have the properties of being stable to the conditions of peptide linkage formation while being readily removable 20 without destruction of the growing peptide chain or racemization of any of the chiral centers contained therein. Suitable protecting groups are 9fluorenylmethyloxycarbonyl (Fmoc), t-butyloxycarbonyl (Boc), benzyloxycarbonyl (Cbz), biphenylisopropyloxycarbonyl, t-25 amyloxycarbonyl, isobornyloxycarbonyl, α , α -dimethyl-3,5dimethoxybenzyloxycarbonyl, o-nitrophenylsulfenyl, 2-cyano-tbutyloxycarbonyl, and the like. The 9-fluorenyl-methyloxycarbonyl (Fmoc) protecting group is particularly preferred for the synthesis of Itherapeutic peptide fragments. Other preferred side chain protecting groups are, for side chain amino groups like lysine and arginine. 30 2,2,5,7,8-pentamethylchroman-6-sulfonyl (pmc), nitro, p-toluenesulfonyl, 4-methoxybenzene-sulfonyl, Cbz, Boc, and adamantyloxycarbonyl; for

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tyrosine, benzyl, o-bromobenzyloxycarbonyl, 2,6-dichlorobenzyl, isopropyl, t-butyl (t-Bu), cyclohexyl, cyclopenyl and acetyl (Ac); for serine, t-butyl, benzyl and tetrahydropyranyl; for histidine, trityl, benzyl, Cbz, p-toluenesulfonyl and 2,4-dinitrophenyl; for tryptophan, formyl; for asparticacid and glutamic acid, benzyl and t-butyl and for cysteine, triphenylmethyl (trityl).

In the solid phase peptide synthesis method, the α -C-terminal amino acid is attached to a suitable solid support or resin. Suitable solid supports useful for the above synthesis are those materials which are inert to the reagents and reaction conditions of the stepwise condensation-deprotection reactions, as well as being insoluble in the media used. The preferred solid support for synthesis of α -C-terminal carboxy peptides is 4-hydroxymethylphenoxymethyl-copoly(styrene-1% divinylbenzene). The preferred solid support for α -C-terminal amide peptides is the 4-(2',4'-dimethoxyphenyl-Fmocaminomethyl)phenoxyacetamidoethyl resin available from Applied Biosystems (Foster City, Calif.). The α -C-terminal amino acid is coupled to the resin by means of N,N'-dicyclohexylcarbodiimide (DCC), N,N'diisopropylcarbodiimide (DIC) or O-benzotriazol-1-yl-N,N,N',N'tetramethyluronium-hexafluorophosphate (HBTU), with or without 4dimethylaminopyridine (DMAP), 1-hydroxybenzotriazole (HOBT), benzotriazol-1-yloxy-tris(dimethylamino)phosphoniumhexafluorophosphate (BOP) or bis(2-oxo-3-oxazolidinyl)phosphine chloride (BOPCI), mediated coupling for from about 1 to about 24 hours at a temperature of between 10° and 50°C. in a solvent such as dichloromethane or DMF.

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When the solid support is 4-(2',4'-dimethoxyphenyl-Fmocaminomethyl)phenoxy-acetamidoethyl resin, the Fmoc group is cleaved with a secondary amine, preferably piperidine, prior to coupling with the α -C-terminal amino acid as described above. The preferred method for coupling to the deprotected 4-(2',4'-dimethoxyphenyl-Fmocaminomethyl)phenoxy-acetamidoethyl resin is O-benzotriazol-1-yl-

N,N,N',N'-tetramethyluroniumhexafluoro-phosphate (HBTU, 1 equiv.) and 1-hydroxybenzotriazole (HOBT, 1 equiv.) in DMF. The coupling of successive protected amino acids can be carried out in an automatic polypeptide synthesizer as is well known in the art. In a preferred embodiment, the α-N-terminal amino acids of the growing peptide chain are protected with Fmoc. The removal of the Fmoc protecting group from the α-N-terminal side of the growing peptide is accomplished by treatment with a secondary amine, preferably piperidine. Each protected amino acid is then introduced in about 3-fold molar excess, and the coupling is preferably carried out in DMF. The coupling agent is normally O-benzotriazol-1-yl-N,N,N',N'-tetramethyluroniumhexafluorophosphate (HBTU, 1 equiv.) and 1-hydroxybenzotriazole (HOBT, 1 equiv.).

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At the end of the solid phase synthesis, the polypeptide is removed from the resin and deprotected, either in successively or in a single operation. Removal of the polypeptide and deprotection can be accomplished in a single operation by treating the resin-bound polypeptide with a cleavage reagent comprising thianisole, water, ethanedithiol and trifluoroacetic acid. In cases wherein the $\alpha\text{-C-terminal}$ of the polypeptide is an alkylamide, the resin is cleaved by aminolysis with an alkylamine. Alternatively, the peptide may be removed by transesterification, e.g. with methanol, followed by aminolysis or by direct transamidation. The protected peptide may be purified at this point or taken to the next step directly. The removal of the side chain protecting groups is accomplished using the cleavage cocktail described above. The fully deprotected peptide is purified by a sequence of chromatographic steps employing any or all of the following types: ion exchange on a weakly basic resin (acetate form); hydrophobic adsorption chromatography on underivitized polystyrene-divinylbenzene (for example, Amberlite XAD); silica gel adsorption chromatography; ion exchange chromatography on carboxymethylcellulose; partition chromatography, e.g. on Sephadex G-25, LH-20 or countercurrent distribution; high performance liquid chromatography (HPLC), especially

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reverse-phase HPLC on octyl- or octadecylsilyl-silica bonded phase column packing.

Molecular weights of these therapeutic peptides are determined using Fast Atom Bombardment (FAB) Mass Spectroscopy.

The therapeutic peptides of the invention may be synthesized with N- and C-terminal protecting groups for use as pro-drugs.

(1) **N-Terminal Protective Groups**

As discussed above, the term "N-protecting group" refers to those groups intended to protect the $\alpha\text{-N-terminal}$ of an amino acid or peptide 10 or to otherwise protect the amino group of an amino acid or peptide against undesirable reactions during synthetic procedures. Commonly used N-protecting groups are disclosed in Greene, "Protective Groups In Organic Synthesis." (John Wiley & Sons, New York (1981)), which is hereby incorporated by reference. Additionally, protecting groups can 15 be used as pro-drugs which are readily cleaved in vivo, for example, by enzymatic hydrolysis, to release the biologically active parent. α-Nprotecting groups comprise loweralkanovl groups such as formyl, acetyl ("Ac"), propionyl, pivaloyl, t-butylacetyl and the like; other acyl groups include 2-chloroacetyl, 2-bromoacetyl, trifluoroacetyl, trichloroacetyl, 20 phthalyl, o-nitrophenoxyacetyl, -chlorobutyryl, benzoyl, 4-chlorobenzoyl, 4-bromobenzoyl, 4-nitrobenzoyl and the like; sulfonyl groups such as benzenesulfonyl, p-toluenesulfonyl and the like; carbamate forming groups such as benzyloxycarbonyl, p-chlorobenzyloxycarbonyl, pmethoxybenzyloxycarbonyl, p-nitrobenzyloxycarbonyl, 2nitrobenzyloxycarbonyl, p-bromobenzyloxycarbonyl, 3,4dimethoxybenzyloxycarbonyl, 3,5-dimethoxybenzyloxycarbonyl, 2,4dimethoxybenzyloxycarbonyl, 4-ethoxybenzyloxycarbonyl, 2-nitro-4,5dimethoxybenzyloxycarbonyl, 3,4,5-trimethoxybenzyloxycarbonyl, 1-(pbiphenylyl)-1-methylethoxycarbonyl, α , α -dimethyl-3,5-30 dimethoxybenzyloxycarbonyl, benzhydryloxycarbonyl, tbutyloxycarbonyl, diisopropylmethoxycarbonyl, isopropyloxycarbonyl,

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ethoxycarbonyl, methoxycarbonyl, allyloxycarbonyl, 2,2,2,trichloroethoxycarbonyl, phenoxycarbonyl, 4-nitrophenoxycarbonyl, fluorenyl-9-methoxycarbonyl, cyclopentyloxycarbonyl, adamantyloxycarbonyl, cyclohexyloxycarbonyl, phenylthiocarbonyl and the like; arylalkyl groups such as benzyl, triphenylmethyl, benzyloxymethyl, 9-fluorenylmethyloxycarbonyl (Fmoc) and the like and silyl groups such as trimethylsilyl and the like.

Carboxy Protective Groups (2)

As discussed above, the term "carboxy protecting group" refers to a carboxylic acid protecting ester or amide group employed to block or protect the carboxylic acid functionality while the reactions involving other functional sites of the compound are performed. Carboxy protecting groups are disclosed in Greene, "Protective Groups in Organic Synthesis" pp. 152-186 (1981), which is hereby incorporated by reference. Additionally, a carboxy protecting group can be used as a pro-drug whereby the carboxy protecting group can be readily cleaved in vivo, for example by enzymatic hydrolysis, to release the biologically active parent. Such carboxy protecting groups are well known to those skilled in the art, having been extensively used in the protection of carboxyl groups in the penicillin and cephalosporin fields as described in U.S. Pat. Nos. 3,840,556 and 3,719,667, the disclosures of which are hereby incorporated herein by reference. Representative carboxy protecting groups are C₁ -C₈ loweralkyl (e.g., methyl, ethyl or t-butyl and the like); arylalkyl such as phenethyl or benzyl and substituted derivatives thereof such as alkoxybenzyl or nitrobenzyl groups and the like: arylalkenyl such as phenylethenyl and the like; aryl and substituted derivatives thereofsuch as 5-indanyl and the like; dialkylaminoalkyl such as dimethylaminoethyl and the like); alkanoyloxyalkyl groups such as acetoxymethyl, butyryloxymethyl, valeryloxymethyl, isobutyryloxymethyl, isovaleryloxymethyl, 1-(propionyloxy)-1-ethyl, 1-(pivaloyloxyl)-1-ethyl, 1methyl-1-(propionyloxy)-1-ethyl, pivaloyloxymethyl, propionyloxymethyl

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and the like; cycloalkanoyloxyalkyl groups such as cyclopropylcarbonyloxymethyl, cyclobutylcarbonyloxymethyl, cyclopentylcarbonyloxymethyl, cyclohexylcarbonyloxymethyl and the like; aroyloxyalkyl such as benzoyloxymethyl, benzoyloxyethyl and the like; arylalkylcarbonyloxyalkyl such as benzylcarbonyloxymethyl, 2benzylcarbonyloxyethyl and the like; alkoxycarbonylalkyl or cycloalkyloxycarbonylalkyl such as methoxycarbonylmethyl, cyclohexyloxycarbonylmethyl, 1-methoxycarbonyl-1-ethyl and the like; alkoxycarbonyloxyalkyl or cycloalkyloxycarbonyloxyalkyl such as methoxycarbonyloxymethyl, t-butyloxycarbonyloxymethyl, 1-10 ethoxycarbonyloxy-1-ethyl, 1-cyclohexyloxycarbonyloxy-1-ethyl and the like: aryloxycarbonyloxyalkyl such as 2-(phenoxycarbonyloxy)ethyl, 2-(5indanyloxycarbonyloxy)ethyl and the like; alkoxyalkylcarbonyloxyalkyl such as 2-(1-methoxy-2-methylpropan-2-oyloxy)ethyl and like; arvialkyloxycarbonyloxyalkyl such as 2-(benzyloxycarbonyloxy)ethyl and 15 the like; arylalkenyloxycarbonyloxyalkyl such as 2-(3-phenylpropen-2yloxycarbonyloxy)ethyl and the like; alkoxycarbonylaminoalkyl such as tbutyloxycarbonylaminomethyl and the like; alkylaminocarbonylaminoalkyl such as methylaminocarbonylaminomethyl and the like; alkanoylaminoalkyl such 20 as acetylaminomethyl and the like; heterocycliccarbonyloxyalkyl such as 4-methylpiperazinylcarbonyloxymethyl and the like; dialkylaminocarbonylalkyl such as dimethylaminocarbonylmethyl, diethylaminocarbonylmethyl and the like; (5-(loweralkyl)-2-oxo-1,3dioxolen-4-yl)alkyl such as (5-t-butyl-2-oxo-1,3-dioxolen-4-yl)methyl and 25 the like; and (5-phenyl-2-oxo-1,3-dioxolen-4-yl)alkyl such as (5-phenyl-2-

Representative amide carboxy protecting groups are aminocarbonyl and loweralkylaminocarbonyl groups.

oxo-1,3-dioxolen-4-yl)methyl and the like.

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Preferred carboxy-protected compounds of the invention are compounds wherein the protected carboxy group is a loweralkyl, cycloalkyl or arylalkyl ester, for example, methyl ester, ethyl ester, propyl ester, isopropyl ester, butyl ester, sec-butyl ester, isobutyl ester, amyl ester, isoamyl ester, octyl ester, cyclohexyl ester, phenylethyl ester and the like or an alkanoyloxyalkyl, cycloalkanoyloxyalkyl, aroyloxyalkyl or an arylalkylcarbonyloxyalkyl ester. Preferred amide carboxy protecting groups are loweralkylaminocarbonyl groups. For example, aspartic acid may be protected at the α -C-terminal by an acid labile group (e.g. tbutyl) and protected at the β -C-terminal by a hydrogenation labile group (e.g. benzyl) then deprotected selectively during synthesis.

Alternatively, it is also possible to obtain fragments of the peptides by fragmenting the naturally occurring amino acid sequence, using, for example, a proteolytic enzyme according to methods well known in the art. Further, it is possible to obtain the desired fragments of the therapeutic peptide through the use of recombinant DNA technology using methods well known in the art.

Modification of Therapeutic Peptides 20 4.

The manner of producing the modified therapeutic peptides of the present invention will vary widely, depending upon the nature of the various elements comprising the molecule. The synthetic procedures will be selected so as to be simple, provide for high yields, and allow for a highly purified stable product. Normally, the reactive group will be created as the last stage, for example, with a carboxyl group, esterification to form an active ester will be the last step of the synthesis. Specific methods for the production of modified therapeutic peptides of the present invention are described below.

Generally, the modified therapeutic peptides of the present invention may be made using blind or structure activity relationship (SAR) driven substitution. SAR is an analysis which defines the

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relationship between the structure of a molecule and its pharmacological activity for a series of compounds. Various studies relative to individual therapeutic peptides show how the activity of the peptide varies according to the variation of chemical structure or chemical properties. More specifically, first the therapeutic activity of the free peptide is

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More specifically, first the therapeutic activity of the free peptide is assayed. Next, the peptide is modified according to the invention, either at the N-terminus, at the C-terminus, or in the interior of the peptide with the linking group only. The linking group will include the reactive group as discussed above. The therapeutic activity of this modified peptide-linking group is assayed next, and based on the detected activity a decision is made regarding the modification site. Next, the peptide

decision is made regarding the modification site. Next, the peptide conjugate is prepared and its therapeutic ativity is determined. If the therapeutic activity of the peptide after conjugation is not substantially reduced (i.e. if the therapeutic activity is reduced by less than 10 fold),

then the stability of the peptide is measured as indicated by its *in vivo* lifetime. If the stability is not improved to a desired level, then the peptide is modified at an alternative site, and the procedure is repeated until a desired level of therapeutic activity and a desired stability are achieved.

More specifically, each therapeutic peptide selected to undergo the derivatization with a linker and a reactive group will be modified according to the following criteria: if a terminal carboxylic group is available on the therapeutic peptide and is not critical for the retention of pharmacological activity, and no other sensitive functional group is present on the therapeutic peptide, then the carboxylic acid will be chosen as attachment point for the linker-reactive group modification. If the terminal carboxylic group *is* involved in pharmacological activity, or if no carboxylic acids are available, then any other sensitive functional group not critical for the retention of pharmacological activity will be selected as the attachment point for the linker-reactive group modification. If several sensitive functional groups are available on a therapeutic peptide, a combination of protecting groups will be used in

such a way that after addition of the linker/reactive group and deprotection of all the protected sensitive functional groups, retention of pharmacological activity is still obtained. If no sensitive functional groups are available on the therapeutic peptide, or if a simpler modification route is desired, synthetic efforts will allow for a modification of the original peptide in such a way that retention of biological activity and retention of receptor or target specificity is obtained. In this case the modification will occur at the opposite end of the peptide.

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An NHS derivative may be synthesized from a carboxylic acid in absence of other sensitive functional groups in the therapeutic peptide. Specifically, such a therapeutic peptide is reacted with N-hydroxysuccinimide in anhydrous CH₂ Cl₂ and EDC, and the product is purified by chromatography or recrystallized from the appropriate solvent system to give the NHS derivative.

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Alternatively, an NHS derivative may be synthesized from a therapeutic peptide that contains an amino and/or thiol group and a carboxylic acid. When a free amino or thiol group is present in the molecule, it is preferable to protect these sensitive functional groups prior to perform the addition of the NHS derivative. For instance, if the molecule contains a free amino group, a transformation of the amine into a Fmoc or preferably into a tBoc protected amine is necessary prior to perform the chemistry described above. The amine functionality will not be deprotected after preparation of the NHS derivative. Therefore this method applies only to a compound whose amine group is not required to be freed to induce a pharmacological desired effect. If the amino group needs to be freed to retain the original biological properties of the molecule, then another type of chemistry described in example 3-6 has to be performed.

In addition, an NHS derivative may be synthesized from a therapeutic peptide containing an amino or a thiol group and no carboxylic acid. When the selected molecule contains no carboxylic acid, an array of bifunctional linkers can be used to convert the molecule

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into a reactive NHS derivative. For instance, ethylene glycolbis(succinimydylsuccinate) (EGS) and triethylamine dissolved in DMF and added to the free amino containing molecule (with a ratio of 10:1 in favor of EGS) will produce the mono NHS derivative. To produce an NHS derivative from a thiol derivatized molecule, one can use N-[- maleimidobutyryloxy]succinimide ester (GMBS) and triethylamine in DMF. The maleimido group will react with the free thiol and the NHS derivative will be purified from the reaction mixture by chromatography on silica or by HPLC.

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An NHS derivative may also be synthesized from a therapeutic peptide containing multiple sensitive functional groups. Each case will have to be analyzed and solved in a different manner. However, thanks to the large array of protecting groups and bifunctional linkers that are commercially available, this invention is applicable to any therapeutic peptide with preferably one chemical step only to derivatize the therapeutic peptide (as described in example 1 or 3) or two steps (as described in example 2 and involving prior protection of a sensitive group) or three steps (protection, activation and deprotection). Under exceptional circumstances only, would we require to use multiple steps (beyond three steps) synthesis to transform a therapeutic peptide into an active NHS or maleimide derivative.

A maleimide derivative may also be synthesized from a therapeutic peptide containing a free amino group and a free carboxylic acid. To produce a maleimide derivative from a amino derivatized molecule, one can use N-[γ-m a leimidobutyryloxy]succinimide ester (GMBS) and triethylamine in DMF. The succinimide ester group will react with the free amino and the maleimide derivative will be purified from the reaction mixture by crystallization or by chromatography on silica or by HPLC.

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Finally, a maleimide derivative may be synthesized from a therapeutic peptide containing multiple other sensitive functional groups and no free carboxylic acids. When the selected molecule contains no carboxylic acid, an array of bifunctional crosslinking reagents can be used to convert the molecule into a reactive NHS derivative. For instance maleimidopropionic acid (MPA) can be coupled to the free amine to produce a maleimide derivative through reaction of the free amine with the carboxylic group of MPA using HBTU/HOBt/DIEA activation in DMF. Alternatively, a lysine residue can be added on the C-terminus end of the peptide to allow for conjugation onto the —amino group of the lysine as described in the examples below. This added lysine allows for simple and efficient modification at the C-terminus of the peptide while keeping the terminal end capped by an amide function as designed by the initial choice of the resin

Many other commercially available heterobifunctional crosslinking reagents can alternatively be used when needed. A large number of bifunctional compounds are available for linking to entities. Illustrative reagents include: azidobenzoyl hydrazide, N-[4-(p-azidosalicylamino)butyl]-3'-[2'-pyridyldithio)propionamide), bissulfosuccinimidyl suberate, dimethyl adipimidate, disuccinimidyl tartrate, N-y-maleimidobutyryloxysuccinimide ester, N-hydroxy sulfosuccinimidyl-4-azidobenzoate, N-succinimidyl [4-azidophenyl]-1,3'-dithiopropionate, N-succinimidyl [4-iodoacetyl]aminobenzoate, glutaraldehyde, and succinimidyl 4-[N-maleimidomethyl]cyclohexane-1-carboxylate.

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Even more specifically, the peptides are preferably modified according to the nature of their substituants and the presence or absence of free cysteines. Most peptides can be gathered into three distinct categories: (1) peptides that contain no cysteines; (2) peptides that contain one cysteine, (3) peptides that contain two cysteines as a disulfide bridge (cystine); and (4) peptides that contain multiple cysteines.

A. Peptides that Contain No Cysteines

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Where the peptide contains no cysteine, addition from the C terminus is performed with all residues cleaved from the support resin and fully protected. Solution phase activation of C-terminus with EDC and NHS can be reacted with an amino-alkyl-maleimide in one pot. The peptide is then fully deprotected. Alternatively, a lysine residue can be added on the C-terminus of the peptide to allow modification at the epsilon amino group of the lysine while keeping the carboxy terminus capped with an amide group. Such an addition of a lysine residue is preferably performed only where the addition does not substantially affect the therapeutic activity of the peptide. The generalized reaction scheme for C-terminus modification of peptides that contain no cysteines is illustrated in the schematic diagram below.

If an N-terminus modification is favored, and again for a peptide containing no cysteine, addition on the N terminus is performed with all residues still on the support resin and fully protected. Addition of activated NHS-Mal bifunctional linker could be performed on deprotected N-terminus with peptide still on resin. The peptide is then fully deprotected. Examples of therapeutic peptides that contain no cystein and undergo a C-terminus modification are described in examples 7-26.

Examples of therapeutic peptides that contain no cystein and undergo a N-terminus modification are described in examples 27-38. The generalized reaction scheme for N-terminus modification of peptides that contain no cysteines is illustrated in the schematic diagrams below, using hetero NHS maleimide (GMBS like) and 3-MPA, respectively.

$$H_2N$$
 Aaa
 CO_2R
 $R = NH_2 \text{ or } H$
 $Aaa = \text{ an amino acid}$
 $Aaa = \text{ alkyl or aromatic}$
 $Aaa = \text{ alkyl or aromatic}$

Hetero NHS Maleimide (GMBS like)

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3-MPA

Alternatively, the peptide may be modified at an internal amino acid (i.e. neither at the C-terminus nor at the N-terminus). The

5 generalized reaction scheme for modification at an internal amino acid of a peptide that contains no free cysteines is illustrated in the schematic diagrams below, using homo bis NHS and hetero NHS maleimide.

- 78 **-**

5 Homo bis NHS

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Ac Aaa
$$^{\prime}$$
 $^{\prime}$ $^{\prime}$

Hetero NHS Maleimide (GMBS like)

Peptides that contain no cysteine and can be modified as described above include fragments of the Kringle 5 peptide, of the GLP-1 peptide, of dynorphin A, human growth hormone releasing factor, the 1-24 fragment of human neuropeptide Y, and human secretin. Full description of the chemistry for each of these peptides is reported in the Example section.

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Peptides that Contain One Cysteine B.

Where the peptide contains one cysteine, the cysteine must stay capped after addition of the maleimide. If the cysteine is involved in binding site, assessment has to be made of how much potency is lost is cysteine is capped by a protecting group. If the cysteine can stay capped, then the synthetic path is similar to that described in section A above for either a C or an N terminus modification.

Alternatively, the peptide may be modified at an internal amino acid (i.e. neither at the C-terminus nor at the N-terminus). The generalized reaction scheme for modification at an internal amino acid of a peptide that contains no cysteines is illustrated in the schematic diagram below, using homobis maleimide and hetero NHS maleimide (GMBS like).

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Homobis Maleimide

Hetero NHS Maleimide (GMBS like)

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Examples of therapeutic peptides that contain one cysteine include G_{α} (the alpha subunit of Gtherapeutic peptide binding protein), the 724-739 fragment of rat brain nitric oxide synthase blocking peptide, the alpha subunit 1-32 fragment of human [Tyr0] inhibin, the 254-274 fragment of HIV envelope protein, and P34cdc2 kinase fragment.

C. Peptides that Contain Two Cysteines as a Disulfide Bridge (Cystine)

15 Where the peptide contains two cysteines as a disulfide bridge, the peptide is cleaved from the support resin before addition of the maleimide. For a modification of the peptide from the C terminus end, all protecting groups are present except at the carboxy terminus (which

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stays unprotected due to cleavage from the support resin) and at the two cysteines, which need to be deprotected when peptide is cleaved from resin. Mild air oxidation yields the disulfide bridge, and the peptide can be purified at that stage. Solution phase chemistry is then required to activate the C-terminus in presence of the disulfide bridge and add the maleimide (through an amino-alkyl-maleimide) to the C-terminus. The peptide is then fully deprotected.

For a modification of the peptide at the N-terminus, the peptide can remain on the support resin. The two cysteines are selectively deprotected before addition of the maleimide. Air oxidation, potentially helped by a catalyst (heterogeneous) can yield the disulfide with the peptide still on the resin. Maleimide is then added on the N-terminus and peptide cleaved from resin and fully deprotected. The generalized reaction scheme for modification at an internal amino acid of a peptide that contains two cysteines in a disulfide bridge is illustrated in the schematic diagram below.

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Alternatively, the peptide may be modified at an internal amino acid (i.e. neither at the C-terminus nor at the N-terminus). The generalized reaction scheme for modification at an internal amino acid of a peptide that contains two cysteines in a disulfide bridge is illustrated in the schematic diagram below.

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Examples of therapeutic peptides that contain two cysteines as a disulfide bridge include human osteocalcin 1-49, human diabetes associated peptide, the 5-28 fragment of human/canine atrial natriuretic peptide, bovine bactenecin, and human [Tyr0]-cortistatin 29.

D. Peptides Containing Multiple Cysteines

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Where the peptide contains multiple cysteines as disulfide

10 bridges, the peptide is cleaved from the support resin before addition of

the maleimide. For a modification of the peptide from the C terminus end, all protecting groups are present except at the carboxy terminus (which stays unprotected due to cleavage from the support resin) and at the two cysteines that are supposed to build a disulfide bridge.

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Cysteines that are involved in other disulfide bridges are deprotected sequencially in pairs using a choice of protecting groups. It is recommended to build and purify each bridge one at a time prior to moving on to the next bridge. Mild air oxidation yields the disulfide bridge, and the peptide should be purified at each stage. Solution phase chemistry is then required to activate the C-terminus in presence of the disulfide bridge and add the maleimide (through an amino-alkylmaleimide) to the C-terminus. The peptide is then fully deprotected.

For a modification of the peptide from the N terminus end, one can leave the peptide on the support resin and selectively deprotect the first two cysteines to build the disulfide under mild air oxidation. Subsequent deprotection will offer the other disulfides before addition of the maleimide. Air oxidation, potentially helped by a catalyst (heterogeneous) can yield the disulfides with the peptide still on the resin. Maleimide is then added on the N-terminus and peptide cleaved from resin and fully deprotected.

Alternatively, the peptide may be modified at an internal amino acid (i.e. neither at the C-terminus nor at the N-terminus).

Peptides containing multiple cysteines include human endothelins and [Lys4] Sarafotoxin S6c.

5. Administration of the Modified Therapeutic Peptides

The modified therapeutic peptide will be administered in a physiologically acceptable medium, e.g. deionized water, phosphate buffered saline (PBS), saline, aqueous ethanol or other alcohol, plasma, proteinaceous solutions, mannitol, aqueous glucose, alcohol, vegetable oil, or the like. Other additives which may be included include buffers, where the media are generally buffered at a pH in the range of about 5

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to 10, where the buffer will generally range in concentration from about 50 to 250 mM, salt, where the concentration of salt will generally range from about 5 to 500 mM, physiologically acceptable stabilizers, and the like. The compositions may be lyophilized for convenient storage and transport.

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The modified Itherapeutic peptides will for the most part be administered orally, parenterally, such as intravascularly (IV), intraarterially (IA), intramuscularly (IM), subcutaneously (SC), or the like. Administration may in appropriate situations be by transfusion. In some instances, where reaction of the functional group is relatively slow, administration may be oral, nasal, rectal, transdermal or aerosol, where the nature of the conjugate allows for transfer to the vascular system. Usually a single injection will be employed although more than one injection may be used, if desired. The modified therapeutic peptides may be administered by any convenient means, including syringe, trocar, catheter, or the like. The particular manner of administration will vary depending upon the amount to be administered, whether a single bolus or continuous administration, or the like. Preferably, the administration will be intravascularly, where the site of introduction is not critical to this invention, preferably at a site where there is rapid blood flow, e.g., intravenously, peripheral or central vein. Other routes may find use where the administration is coupled with slow release techniques or a protective matrix. The intent is that the Itherapeutic peptides be effectively distributed in the blood, so as to be able to react with the blood components. The concentration of the conjugate will vary widely, generally ranging from about 1 pg/ml to 50 mg/ml. The total administered intravascularly will generally be in the range of about 0.1 mg/ml to about 10 mg/ml, more usually about 1 mg/ml to about 5 mg/ml.

By bonding to long-lived components of the blood, such as immunoglobulin, serum albumin, red blood cells and platelets, a number of advantages ensue. The activity of the modified therapeutic peptides compound is extended for days to weeks. Only one administration need

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be given during this period of time. Greater specificity can be achieved, since the active compound will be primarily bound to large molecules, where it is less likely to be taken up intracellularly to interfere with other physiological processes.

The formation of the covalent bond between the blood component may occur *in vivo* or *ex vivo*. For *ex vivo* covalent bond formation, the modified Itherapeutic peptide is added to blood, serum or saline solution containing human serum albumin or IgG to permit covalent bond formation between the modified therapeutic peptide and the blood component. In a preferred format, the therapeutic peptide is modified with maleimide and it is reacted with human serum albumin in saline solution. Once the modified therapeutic peptide has reacted with the blood component, to form a therapeutic peptide-protein conjugate, the conjugate may be administered to the patient.

Alternatively, the modified therapeutic peptide may be administered to the patient directly so that the covalent bond forms between the modified Itherapeutic peptide and the blood component *in vivo*.

In addition, where localized delivery of therapeutic peptides is desired, several methods of delivery may be used:

A. Open Surgical Field Lavage

There are a number of indications for local therapeutic compounds which would entail administration of the therapeutic compound as an adjunct to open surgery. In these cases, the therapeutic compound would either be lavaged in the surgical site (or a portion of that site) prior to closure, or the therapeutic compound would be incubated for a short time in a confined space (e.g., the interior of a section of an artery following an endarterectomy procedure or a portion of GI tract during resection) and the excess fluid subsequently evacuated.

B. Incubation of Tissue Grafts

Tissue grafts such as autologous and xenobiotic vein/artery and valve grafts as well as organ grafts can be pretreated with therapeutic

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compounds that have been modified to permit covalent bond formation by either incubating them in a therapeutic solution and/or perfusing them with such a solution.

C. Catheter Delivery

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A catheter is used to deliver the therapeutic compound either as part of an endoscopic procedure into the interior of an organ (e.g., bladder, GI tract, vagina/uterus) or adjunctive to a cardiovascular catheter procedure such as a balloon angioplasty. Standard catheters as well as newer drug delivery and iontophoretic catheters can be utilized.

D. Direct Injection

For certain poorly vascularized spaces such as intra-articular joint spaces, a direct injection of a therapeutic compound may be able to bioconjugate to surface tissues and achieve a desirable duration of drug effect. Other applications could include intra medullary, intratumor, intravaginal, intrauterine, intra intestinal, intra eustachian tube, intrathecal, subcutaneous, intrarticular, intraperitoneal or intraocular injections as weel as via bronchoscope, via nasogastiric tube and via nophrostomy.

6. Monitoring the Presence of Modified Therapeutic Peptide Derivatives

Another aspect of this invention relates to methods for determining the concentration of the therapeutic peptides and/or analogs, or their derivatives and conjugates in biological samples (such as blood) and determining the peptidase stability of the modified peptides. The blood of the mammalian host may be monitored for the presence of the modified therapeutic peptide compounds one or more times. By taking a portion or sample of the blood of the host, one may determine whether the therapeutic peptide has become bound to the long-lived blood components in sufficient amount to be therapeutically active and, thereafter, the level of therapeutic peptide compound in the blood. If desired, one may also determine to which of the blood components the therapeutic peptide derivative molecule is bound. This

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is particularly important when using non-specific therapeutic peptides. For specific maleimide-therapeutic peptides, it is much simpler to calculate the half life of serum albumin and IgG.

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One method for determining the concentration of the therapeutic peptide, analogs, derivatives and conjugates is to use antibodies specific to the therapeutic peptides or therapeutic peptide analogs or their derivatives and conjugates, and to use such antibodies as a treatment for toxicity potentially associated with such therapeutic peptides, analogs, and/or their derivatives or conjugates. This is advantageous because the increased stability and life of the therapeutic peptides *in vivo* in the patient might lead to novel problems during treatment, including increased possibility for toxicity. It should be mentioned, however, that in some cases, the traditional antibody assay may not recognize the difference between cleaved and uncleaved therapeutic peptides. In such cases, other assay techniques may be employed, for example LC/MS (Liquid Chromatography / Mass Spectrometry).

The use of antibodies, either monoclonal or polyclonal, having specificity for a particular therapeutic peptide, analog or derivative thereof, can assist in mediating any such problem. The antibody may be generated or derived from a host immunized with the particular therapeutic peptide, analog or derivative thereof, or with an immunogenic fragment of the agent, or a synthesized immunogen corresponding to an antigenic determinant of the agent. Preferred antibodies will have high specificity and affinity for native, derivatized and conjugated forms of the therapeutic peptide or therapeutic peptide analog. Such antibodies can also be labeled with enzymes, fluorochromes, or radiolabels.

Antibodies specific for derivatized therapeutic peptides may be produced by using purified therapeutic peptides for the induction of derivatized therapeutic peptide-specific antibodies. By induction of antibodies, it is intended not only the stimulation of an immune response by injection into animals, but analogous steps in the production of

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synthetic antibodies or other specific binding molecules such as screening of recombinant immunoglobulin libraries. Both monoclonal and polyclonal antibodies can be produced by procedures well known in the art. In some cases, the use of monoclonal antibodies may be preferred over polyclonal antibodies, such as when degradation occurs over an area not covered by epitope/antibody recognition.

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The antibodies may be used to treat toxicity induced by administration of the therapeutic peptide, analog or derivative thereof, and may be used *ex vivo* or *in vivo*. *Ex vivo* methods would include immuno-dialysis treatment for toxicity employing antibodies fixed to solid supports. *In vivo* methods include administration of antibodies in amounts effective to induce clearance of antibody-agent complexes.

The antibodies may be used to remove the therapeutic peptides, analogs or derivatives thereof, and conjugates thereof, from a patient's blood ex vivo by contacting the blood with the antibodies under sterile conditions. For example, the antibodies can be fixed or otherwise immobilized on a column matrix and the patient's blood can be removed from the patient and passed over the matrix. The therapeutic peptide analogs, derivatives or conjugates, will bind to the antibodies and the blood containing a low concentration of the therapeutic peptide, analog, derivative or conjugate, then may be returned to the patient's circulatory system. The amount of therapeutic peptide compound removed can be controlled by adjusting the pressure and flow rate. Preferential removal of the therapeutic peptides, analogs, derivatives and conjugates from the plasma component of a patient's blood can be affected, for example, by the use of a semipermeable membrane, or by otherwise first separating the plasma component from the cellular component by ways known in the art prior to passing the plasma component over a matrix containing the anti-therapeutic antibodies. Alternatively the preferential removal of therapeutic peptide-conjugated blood cells, including red blood cells, can be effected by collecting and concentrating the blood cells in the patient's blood and contacting those cells with fixed anti-therapeutic

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antibodies to the exclusion of the serum component of the patient's blood.

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The antibodies can be administered *in vivo*, parenterally, to a patient that has received the therapeutic peptide, analogs, derivatives or conjugates for treatment. The antibodies will bind the therapeutic peptide compounds and conjugates. Once bound the therapeutic peptide, activity will be hindered if not completely blocked thereby reducing the biologically effective concentration of therapeutic peptide compound in the patient's bloodstream and minimizing harmful side effects. In addition, the bound antibody–therapeutic peptide complex will facilitate clearance of the therapeutic peptide compounds and conjugates from the patient's blood stream.

The invention having been fully described is now exemplified by the following non-limiting examples.

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EXAMPLES

A. General Method of Synthesis of a Modified Therapeutic Peptide

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Solid phase peptide synthesis of the modified peptide on a 100 µmole scale was performed on a Symphony Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin, Fmoc protected amino acids, O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) in N,N-dimethylformamide (DMF) solution and activation with N-methyl morpholine (NMM), and piperidine deprotection of Fmoc groups (Step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by either the coupling of 3-maleimidopropionic acid (3-MPA), the coupling of acetic acid or the coupling of one or multiple Fmoc-AEEA followed by the coupling of 3-MPA (Step 3). Resin cleavage and products isolation are performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The products are purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 μm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product should have >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

B. Alteration of the Native Peptide Chain

To facilitate modification of the peptide, one or more amino acid residues may be added to the peptide as described in examples 1 to 5, and/or one or more amino acid residues may be replaced with other amino acid residues. This alteration aids attachment of the reactive group.

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Example 1 — Addition of Lys at C-Terminus of Kringle-5 Preparation of NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂.3TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH. Deblocking of the Fmoc group the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling. Final cleavage from the resin was performed using cleavage mixture as described above. The product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization.

15 <u>Example 2</u> - Addition of Lys at C-Terminus of Kringle-5 Preparation of NAc-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂.2TFA.3TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH. Deblocking of the Fmoc group the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling. Final cleavage from the resin was performed using cleavage mixture as described above. The product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization.

30 <u>Example 3</u> - Addition of Lys at N-Terminus of Kringle-5 Preparation of NAc-Tyr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂.3TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH. Deblocking of the Fmoc group the the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling. Final cleavage from the resin was performed using cleavage mixture as described above. The product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization.

15 <u>Example 4</u> - Addition of Lys at N-Terminus of Kringle-5, Substitution of Cys with Ala at Position 524 Preparation of NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Ala⁵²⁴-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂.4TFA

Using automated peptide synthesis, the following protected amino 20 acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tvr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Ala-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, 25 Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Arg(Pbf)-OH. Deblocking of the Fmoc group the the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under 30 conditions similar to amino acid coupling. Final cleavage from the resin was performed using cleavage mixture as described above. The

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product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization.

Example 5 - Addition of Lys at N-Terminus of Kringle-5 Preparation of NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Lys-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Gly-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Arg(Pbf)-OH. Deblocking of the Fmoc group the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling. Final cleavage from the resin was performed using cleavage mixture as described above. The product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization.

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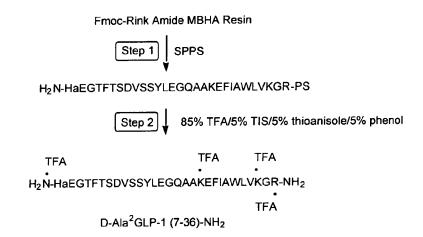
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Example 6 - Preparation of D-Ala2 GLP-1 (7-36) Amide

Solid phase peptide synthesis of the GLP-1 analog on a 100 µmole scale is performed using manual solid-phase synthesis and a Symphony Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin, Fmoc protected amino acids, O-benzotriazol-1-yl-N, N, N, N-tetramethyl-uronium hexafluorophosphate (HBTU) in N,N-dimethylformamide (DMF) solution and activation with N-methyl morpholine (NMM), and piperidine deprotection of Fmoc groups (Step 1). Resin cleavage and product isolation is performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 2). The product is purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and

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0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired peptide in >95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic diagram below.



C. Preparation of Modified Peptides From Peptides Containing No Cysteines

Preparation of maleimido peptides from therapeutic peptides containing multiple protected functional groups and no Cysteine is exemplified by the synthesis of peptides as described below. The peptide may be modified at the N-terminus, the C-terminus, or at an amino acid located between the N-terminus and the C-terminus. The modified peptide is synthesized by linking off the N-terminus of the natural peptide sequence or by linking off the modified C-terminus of the natural peptide sequence. One or more additional amino acids may be added to the therapeutic peptide to facilitate attachment of the reactive group.

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Modification of the Therapeutic Peptide at the C-Terminus

Example 7 – Modification of RSV Peptide at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of Val-Ile-Thr-Ile-Glu-Leu-Ser-Asn-Ile-Lys-Glu-Asn-Lys-Met-Asn-Gly-Ala-Lys-Val-Lys-Leu-Ile-Lys-Gln-Glu-Leu-Asp-Lys-Tyr-Lys-Asn-Ala-Val-Lys-(Νε-ΜΡΑ)

Solid phase peptide synthesis of the DAC analog on a 100 µmole 10 scale is performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Lys(Aloc)-OH, Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asp(tBu)-15 OH, Fmoc-Leu-OH, Fmoc-Glu(tBu)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ile-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Val-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ala-OH, Fmoc-Gly-OH, Fmoc-Asn(Trt)-OH, Fmoc-Met-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ile-OH, Fmoc-Asn(Trt)-20 OH, Fmoc-Ser(tBu)-OH, Fmoc-Leu-OH, Fmoc-Glu(tBu)-OH, Fmoc-Ile-OH, Fmoc-Thr(tBu)-OH, Fmoc-Ile-OH, Fmoc-Val-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). 25 Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh3)4 dissolved in 5 mL of CHCl3:NMM:HOAc (18:1:0.5) for 2 h 30 (Step 2). The resin is then washed with CHCl3 (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with N,N-

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dimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et2O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H2O (A) and 0.045% TFA in CH3CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

Example 8 – Modification of Dyn A 1-13 at the ϵ -Amino Group of the Added C-terminus Lysine Residue - Synthesis of Dyn A 1-13(N ϵ -MPA)-NH₂

15 Tyr-Gly-Gly-Phe-Leu-Arg-Arg-Ile-Arg-Pro-Lys-Leu-Lys-(Nε-MPA)-NH₂

Solid phase peptide synthesis of a modified Dyn A 1-13 on a 100 umole scale was performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. 20 The following protected amino acids were sequentially added to resin: Fmoc-Lys(Aloc)-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Pro-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ile-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Phe-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Tyr(tBu)-OH. They were dissolved in N,N-dimethylformamide 25 (DMF) and, according to the sequence, activated using O-benzotriazol-1-vl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and

dimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5

mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with N, N-dimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et_2O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH_3CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

The structure of this product is

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Example 9 - Modification of Dyn A 2-13 at the ε-Amino Group of the Added C-terminus Lysine Residue - Synthesis of Dyn A 2-13(Nε-MPA)-NH₂ Gly-Gly-Phe-Leu-Arg-Arg-Ile-Arg-Pro-Lys-Leu-Lys-(Nε-MPA)-NH₂

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Mtt)-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Pro-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Phe-OH, Fmoc-Gly-OH, and Boc-Gly-OH. Manual synthesis was employed for the remaining steps: selective removal of the Mtt group and coupling of MPA using HBTU/HOBt/DIEA activation in

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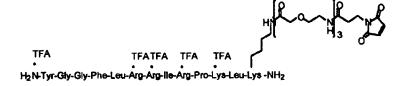
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DMF. The target dynorphin analog was removed from the resin; the product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization in a 35% yield. Anal. HPLC indicated product to be >95% pure with R_t = 30.42 min. ESI-MS m/z for $C_{73}H_{123}N_{25}O_{15}$ (MH⁺), calcd 1590.0, found MH³⁺ 531.3.

Example 10 - Modification of Dyn A 1-13 at the ε-Amino Group of the Added C-terminus Lysine Residue - Synthesis of Dyn A 1-13(AEA₃-MPA)-NH₂ Tyr-Gly-Gly-Phe-Leu-Arg-Arg-Ile-Arg-Pro-Lys-Leu-Lys-(AEA3-MPA)-NH2

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Mtt)-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Pro-OH, Fmoc-15 Arg(Pbf)-OH, Fmoc-Ile-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Phe-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, and Boc-Tyr(Boc)-OH. Manual synthesis was employed for the remaining steps: selective removal of the Mtt group, the coupling of three-Fmoc-AEA-OH groups (AEA = aminoethoxyacetic acid) with Fmoc removal in-between each coupling, and MPA acid using HBTU/HOBt/DIEA activation in DMF. The target dynorphin analog was removed from the resin; the product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization in a 29% yield. Anal. HPLC indicated product to be >95% pure with Rt = 33.06 min. ESI-MS m/z for $C_{94}H_{154}N_{29}O_{23}$ (MH⁺), calcd 2057.2, found MH⁴⁺ 515.4, MH³⁺ 686.9, MH²⁺ 1029.7.



Example 11 – Modification of Dyn A 2-13 at the ε-Amino Group of the Added C-terminus Lysine Residue - Synthesis of Dyn A 2-13(AEA₃-MPA)-NH₂ Gly-Gly-Phe-Leu-Arg-Arg-Ile-Arg-Pro-Lys-Leu-Lys-(AEA₃-MPA)-NH₂

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Mtt)-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Pro-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Phe-OH, Fmoc-Gly-OH, and Fmoc-Gly-OH. Manual synthesis was employed for the remaining steps: selective removal of the Mtt group, the coupling of three-Fmoc-AEA-OH groups, with Fmoc removal in-between each coupling, and MPA using HBTU/HOBt/DIEA activation in DMF. The target dynorphin analog was removed from the resin; the product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization in a 29% yield. Anal. HPLC indicated product to be >95% pure with $R_t = 31.88$ min. ESI-MS m/z for $C_{85}H_{145}N_{25}O_{21}$ (MH $^+$), calcd 1894.3, found MH $^{4+}$ 474.6, MH $^{3+}$ 632.4, MH $^{2+}$ 948.10.

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Example 12 – Modification of Neuropeptide Y at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of Tyr-Pro-Ser-Lys-Pro-Asp-Asn-Pro-Gly-Glu-Asp-Ala-Pro-Ala-Glu-Asp-Met-Ala-Arg-Tyr-Tyr-Ser-Ala-Leu-Lys-(N-εMPA)-NH₂

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Solid phase peptide synthesis of a modified neuropeptide Y analog on a 100 µmole scale is performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink

Amide MBHA. The following protected amino acids were sequentially added to resin: Fmoc-Lys(Aloc)-OH, Fmoc-Leu-OH, Fmoc-Ala-OH, Fmoc-Ser(tBu)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ala-OH, Fmoc-Met-OH, Fmoc-Asp(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Ala-OH, Fmoc-Pro-OH, Fmoc-Ala-OH, Fmoc-5 Asp(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Gly-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Asp(tBu)-OH, Fmoc-Pro-OH, Fmoc-Lys(Boc)-OH. Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Tyr(tBu)-OH, They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-10 uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a 15 solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of the 3maleimidopropionic acid (Step 3). Between every coupling, the resin is 20 washed 3 times with N,N-dimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient 25 elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenylhexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

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Example 13 - Modification of GLP-1 (7-36) at the C-Terminus Arginine Preparation of GLP-1 (7-36)-EDA-MPA

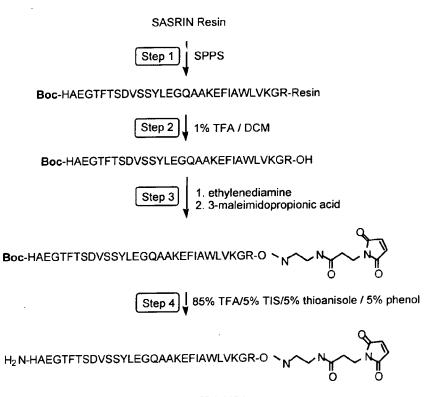
Solid phase peptide synthesis of a modified GLP-1 analog on a 5 100 umole scale is performed manually and on a Symphony Peptide Synthesizer SASRIN (super acid sensitive resin). The following protected amino acids are sequentially added to the resin: Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(Boc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp(Boc)-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-Phe-10 OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ala-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gly-OH, Fmco-Glu(OtBu)-OH, Fmoc-Leu-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Val-OH, Fmoc-Asp(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH. Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-15 Glu(OtBu)-OH, Fmoc-Ala-OH, Boc-His(Trt)-OH. They are dissolved in N.N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The fully protected peptide is cleaved from the resin by treatment with 1% TFA / DCM (Step 2). Ethylenediamine and 3maleimidopropionic acid are then sequentially added to the free Cterminus (Step 3). The protecting groups are then cleaved and the 25 product isolated using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 μm 30 quard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in

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>95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic diagram below.

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GLP-1 (7-36)-EDA-MPA

Example 14 - Modification of Exendin-4 at the C-terminus Serine 10 Preparation of Exendin-4 (1-39)-EDA-MPA

Solid phase peptide synthesis of a modified Exendin-4 analog on a 100 µmole scale is performed manually and on a Symphony Peptide Synthesizer SASRIN (super acid sensitive resin). The following protected amino acids are sequentially added to the resin: Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Cly-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Cly-OH, Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Trp(Boc)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ile-

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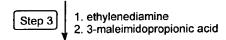
OH, Fmoc-Phe-OH, Fmoc-Leu-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Met-OH, Fmoc-Gln(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Leu-OH, Fmoc-Asp(OtBu)-OH, Fmoc--Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, 5 Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Gly-OH, Boc-His(Trt)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N, Ntetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is 10 achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). The fully protected peptide is cleaved from the resin by treatment with 1% TFA / DCM (Step 2). Ethylenediamine and 3-maleimidopropionic acid are then sequentially added to the free C-terminus (Step 3). The protecting groups are then 15 cleaved and the product isolated using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C_{18} , 60Å, $8~\mu\text{m}$, 21 mm x 25 cm column equipped with a 20 Dynamax C_{18} , 60Å, 8 μm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.





Boc-HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS-Resin

Boc-HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS-OH



Boc-HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS-O

Step 4 85% TFA/5% TIS/5% thioanisole / 5% phenol

Exendin-4 (1-39)-EDA-MPA

Example 15 – Modification of Secretin Peptide at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of His-Ser-Asp-Gly-Thr-Phe-Thr-Ser-Glu-Leu-Ser-Arg-Leu-Arg-Glu-Gly-Ala-Arg-Leu-Glu-Arg-Leu-Gln-Gly-Leu-Val-Lys-(Νε-ΜΡΑ)-NΗ₂

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Solid phase peptide synthesis of a modified secretin peptide analog on a 100 µmole scale is performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Lys(Aloc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, 15 Fmoc-Gly-OH, Fmoc-Gln(Trt)-OH, Fmoc-Leu-OH, Fmoc-Leu-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Leu-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ala-OH, Fmoc-Gly-OH, Fmoc-Glu(tBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Leu-OH, Fmoc-Glu(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-20 OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Asp(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-His(Boc)-OH. They are dissolved in N, Ndimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). 25 Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h 30 (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with N,Ndimethylformamide (DMF) and 3 times with isopropanol. The peptide is 35

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cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

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Example 16 – Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(Nε-MPA)-NH₂.2TFA

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Solid phase peptide synthesis of a modified Kringle-5 peptide on a 100 µmole scale was performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Lys(Aloc)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Asp(tBu)-OH, Fmoc-20 Tyr(tBu)-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH. They were dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is 25 achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (step 1). At the end of the synthesis Acetic Anhydride was added to acetylate the N-terminal. The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ 30 dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 3).

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Between every coupling, the resin is washed 3 times with *N,N*-dimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

Example 17 – Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue
15 Preparation of NAc-Tyr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(Νε-ΜΡΑ)-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-20 Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH (step 1). Deblocking of the Fmoc group the the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Final cleavage from the resin 25 was performed using cleavage mixture as described above. The product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq 30 of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis

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was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

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Example 18 - Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Ala-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(Nε-MPA)-NH₂.3TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lvs(Aloc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, 20 Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Ala-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Gly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Arg(Pbf)-OH (step 1). Deblocking of the Fmoc group the the N-25 terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling. Final cleavage from the resin was performed using cleavage mixture as described above. The product was isolated by precipitation and purified 30 by preparative HPLC to afford the desired product as a white solid upon lyophilization.

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

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Example 19 – Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of NAc-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Pro-Trp-Lys-(Nε-MPA)-NH₂.TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Asp(OtBu)-OH

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followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

Example 20 – Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of NAc-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(Nε-MPA)-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lvs(Aloc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-15 Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH (Step 1). Deblocking of the Fmoc group the N-terminal of the resinbound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling. Final cleavage from the resin 20 was performed using cleavage mixture as described above. The product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 30 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative

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binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH_3CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

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Example 21 - Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Lys-(Nε-MPA)-NH₂.2TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH (Step 1). Deblocking of the Fmoc group the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling. Final cleavage from the resin was performed using cleavage mixture as described above. The product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a

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Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

Example 22 – Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(Nε-AEEA-MPA)-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-10 Lys(Aloc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH (Step 1). Deblocking of the Fmoc group at the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was 15 performed under conditions similar to amino acid coupling. The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc 20 in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition. The synthesis was then reautomated for the addition of the AEEA (aminoethoxyethoxyacetic acid) groupand of the 3-maleimidopropionic acid (MPA) (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% 25 TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 30 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

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Example 23 – Modification of Kringle-5 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of NAc-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-(Νε-ΑΕΕΑ_n-MPA)-NH₂.2TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH (Step 1). Deblocking of the Fmoc group at the N-terminal of the resin-bound amino acid was performed with 20% piperidine in DMF for about 15-20 minutes. Coupling of the acetic acid was performed under conditions similar to amino acid coupling.

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition. The synthesis was then reautomated for the addition of n AEEA (aminoethoxyethoxyacetic acid) groups and of the 3-maleimidopropionic acid (MPA) (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

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Example 24 – Modification of GLP-1 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of GLP-1 (1-36)-Lys 37 (Nε-MPA)-NH₂.5TFA;

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His-Asp-Glu-Phe-Glu-Arg-His-Ala-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Val-Ser-Ser-Tyr-Leu-Glu-Gly-Gln-Ala-Ala-Lys-Glu-Phe-lle-Ala-Trp-Leu-Val-Lys-Gly-Arg-Lys(Nε-MPA)-NH₂.5TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Ala-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ala-OH, Boc-His(N-Trt)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Phe-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Phe-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Phe-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Phe-OH, Fmoc-Phe-OH

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode

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array detector and using electro-spray ionization. These steps are illustrated in the schematic diagram below.

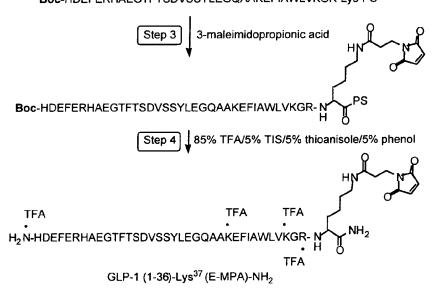
Fmoc-Rink Amide MBHA Resin

Step 1 SPPS

Boc-HDEFERHAEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys(Aloc)-PS

Boc-HDEFERHAEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys-PS

Step 2 JPd(PPh₃)₄/NMM/HOAc/CHCl₃



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Example 25 - Modification of GLP-1 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of GLP-1 (1-36)-Lys 37 (Nε-AEEA-AEEA-MPA)-NH₂.5TFA; His-Asp-Glu-Phe-Glu-Arg-His-Ala-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Val-Ser-Ser-Tyr-Leu-Glu-Gly-Gln-Ala-Ala-Lys-Glu-Phe-Ile-Ala-Trp-Leu-Val-Lys-Gly-Arg-Lys(Nε-AEEA-AEEA-MPA)-NH₂.5TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Ala-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gly-OH, Fmoc-

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Glu(OtBu)-OH, Fmoc-Leu-OH, Fmoc-Tyr(Pbf)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ala-OH, Boc-His(N-Trt)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Asp(OtBu)-OH, Boc-His(N-Trt)-OH (step 1).

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The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the two AEEA (aminoethoxyethoxyacetic acid) groups and the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization, ESI-MS m/z for $C_{174}H_{265}N_{44}O_{56}$ (MH⁺), calcd 3868, found [M+H₂]²⁺ 1934, [M+H₃]³⁺ 1290, [M+H₄]⁴⁺ 967. These steps are illustrated in the schematic diagram below.

Fmoc-Rink Amide MBHA Resin

Boc-HAEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys(Aloc)-PS

Boc-HAEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys-PS

Example 26 - Modification of GLP-1 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of GLP-1 (7-36)-Lys³⁷(Nε-MPA)-NH₂.4TFA; His-Ala-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Val-Ser-Ser-Tyr-Leu-Glu-Gly-Gln-Ala-Ala-Lys-Glu-Phe-Ile-Ala-Trp-Leu-Val-Lys-Gly-Arg-Lys(Nε-MPA)-NH₂.4TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(tBoc)-OH, Fmoc-15 Ala-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Leu-OH, Fmoc-Tyr(Pbf)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH,

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Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ala-OH, Boc-His(N-Trt)-OH (Step 1).

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

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Example 27 - Modification of GLP-1 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of GLP-1 (7-36)-Lys³⁷(Nε-AEEA-AEEA-MPA)-NH₂.4TFA His-Ala-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Val-Ser-Ser-Tyr-Leu-Glu-Gly-Gln-Ala-Ala-Lys-Glu-Phe-Ile-Ala-Trp-Leu-Val-Lys-Gly-Arg-Lys(Nε-AEEA-AEEA-MPA)-NH₂.4TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Leu-OH, Fmoc-Tyr(Pbf)-OH, Fmoc-Ser(tBu)-OH,

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Fmoc-Ser(tBu)-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ala-OH, Boc-His(N-Trt)-OH (Step 1).

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the two AEEA (aminoethoxyethoxyacetic acid) groups and the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

Example 28 - Modification of D-Ala² GLP-1 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of D-Ala² GLP-1 (7-36)-Lys³⁷(Nε-MPA)-NH₂.4TFA His-d-Ala-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Val-Ser-Ser-Tyr-Leu-Glu-Gly-Gln-Ala-Ala-Lys-Glu-Phe-Ile-Ala-Trp-Leu-Val-Lys-Gly-Arg-Lys(Nε-MPA)-NHh₂.4TFA

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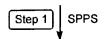
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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Ala-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Leu-OH, Fmoc-Tyr(Pbf)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-d-Ala-OH, Boc-His(N-Trt)-OH (Step1).

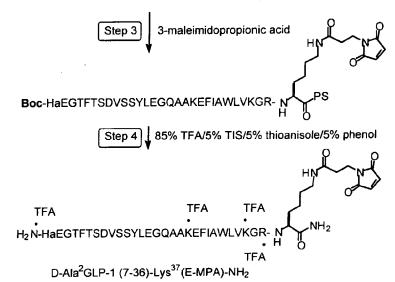
The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization. These steps are illustrated in the schematic diagram below.

Fmoc-Rink Amide MBHA Resin



Boc-HaEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys(Aloc)-PS

Boc-HaEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys-PS



Example 29 - Modification of D-Ala 2 GLP-1 at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of D-Ala 2 GLP-1 (7-36)-Lys 37 (Nε-AEEA-AEEA-MPA)-

Preparation of D-Ala² GLP-1 (7-36)-Lys³ (Nε-AEEA-AEEA-MPA)-NH₂.4TFA
His-D-Ala-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Val-Ser-Ser-Tyr-Leu-Glu-Gly-Gln-Ala-Ala-Lys-Glu-Phe-Ile-Ala-Trp-Leu-Val-Lys-Gly-Arg-Lys (Nε-AEEA-AEEA-MPA)-NH₂.4TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(tBoc)-OH, Fmoc-Ala-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-

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Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-d-Ala-OH, Boc-His(N-Trt)-OH (Step 1).

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq 5 of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the two AEEA (aminoethoxyethoxyacetic acid) groups and the 3-maleimidopropionic 10 acid (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O 15 (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode 20 array detector and using electro-spray ionization. These steps are illustrated in the schematic diagram below.

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Fmoc-Rink Amide MBHA Resin

Boc-HaEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys(Aloc)-PS

Boc-HaEGTFTSDVSSYLEGQAAKEFIAWLVKGR-Lys-PS

Example 30 - Modification of Exendin-4(1-39) at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of Exendin-4 (1-39)-Lys⁴⁰(Nε-MPA)-NH₂; His-Gly-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Leu-Ser-Lys-Gln-Met-Glu-Glu-Glu-Ala-Val-Arg-Leu-Phe-Ile-Glu-Trp-Leu-Lys-Asn-Gly-Gly-Pro-Ser-Ser-Gly-Ala-Pro-Pro-Pro-Ser-Lys (Nε-MPA)-NH₂.5TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ala-OH, Fmoc-Gly-OH, Fmoc-Ser-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Asn(Trt)-OH,
 Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Leu-OH, Fmoc-Arg(Bpf)-OH, Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Leu-OH, Fmoc-Asp(OtBu)-OH,

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Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Boc-His(Trt)-OH (Step 1).

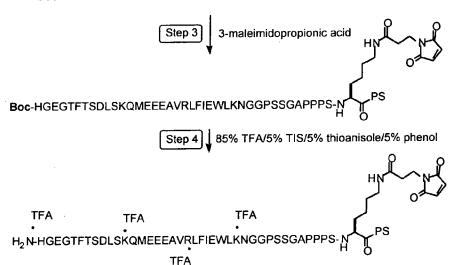
The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq 5 of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 10 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a 15 Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization. These steps are 20 illustrated in the schematic diagram below.

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Fmoc-Rink Amide MBHA Resin

Boc-HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS-Lys(Aloc)-PS

Boc-HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS-Lys-PS



Exendin-4 (1-39)-Lys40 (E-MPA)-NH2

<u>Example 31</u> - Modification of Exendin-4(1-39) at the ϵ -Amino Group of the Added C-terminus Lysine Residue

Preparation of Exendin-4 (1-39)-Lys⁴⁰(Nε-AEEA-AEEA-MPA)-NH₂.5TFA; His-Gly-Glu-Gly-Thr-Phe-Thr-Ser-Asp-Leu-Ser-Lys-Gln-Met-Glu-Glu-Glu-Ala-Val-Arg-Leu-Phe-Ile-Glu-Trp-Leu-Lys-Asn-Gly-Gly-Pro-Ser-Ser-Gly-Ala-Pro-Pro-Pro-Ser-Lys(Nε-AEEA-AEEA-MPA)-NH₂.5TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Ser-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Leu-OH, Fmoc-Arg(Bpf)-OH,

Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Met-OH, Fmoc-Gln(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Leu-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Gly-OH, Boc-His(Trt)-. OH (Step 1).

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The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc 10 in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the two AEEA (aminoethoxyethoxyacetic acid) groups and the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed. using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by 15 precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H2O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV 20 detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization. These steps are illustrated in the schematic diagram below.

Fmoc-Rink Amide MBHA Resin

Boc-HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS-Lys(AlogPS

Boc-HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS-Lys-PS

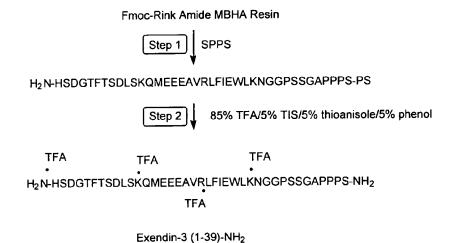
Exendin-4 (1-39)-Lys⁴⁰(E-AEEA₂-MPA)-NH₂

Example 32 - Modification of Exendin-3(1-39) at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of Exendin-3 (1-39)-Lys⁴⁰(Nε-MPA)-NH₂.5TFA; His-Ser-Asp-Gly-Thr-Phe-Thr-Ser-Asp-Leu-Ser-Lys-Gln-Met-Glu-Glu-Glu-Ala-Val-Arg-Leu-Phe-Ile-Glu-Trp-Leu-Lys-Asn-Gly-Gly-Pro-Ser-Ser-Gly-Ala-Pro-Pro-Pro-Ser-Lys(Nε-MPA)-NH₂.5TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ala-OH, Fmoc-Gly-OH, Fmoc-Ser-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Asn(Trt)-OH,
Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Leu-OH, Fmoc-Arg(Bpf)-OH, Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Leu-OH, Fmoc-Asp(OtBu)-OH,

Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(OtBu)-OH, Boc-His(Trt)-OH (Step 1).

The selective deprotection of the Lys(Aloc) group was performed manually and accomplished by treating the resin with a solution of 3 eq 5 of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis was then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation was performed using 10 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a 15 Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization. These steps are 20 illustrated in the schematic diagram below.



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Example 33 - Modification of Exendin-3(1-39) at the ε-Amino Group of the Added C-terminus Lysine Residue Preparation of Exendin-3 (1-39)-Lys⁴⁰(Nε-AEEA-AEEA-MPA)-NH₂.5TFA; His-Ser-Asp-Gly-Thr-Phe-Thr-Ser-Asp-Leu-Ser-Lys-Gln-Met-Glu-Glu-Glu-Ala-Val-Arg-Leu-Phe-Ile-Glu-Trp-Leu-Lys-Asn-Gly-Gly-Pro-Ser-Ser-Gly-Ala-Pro-Pro-Pro-Ser-Lys(Nε-AEEA-AEEA-MPA)-NH₂.5TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Aloc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ala-OH, Fmoc-Gly-OH, Fmoc-Ser-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Trp-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ile-OH, Fmoc-Phe-OH, Fmoc-Leu-OH, Fmoc-Arg(Bpf)-OH, Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Leu-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(OtBu)-OH, Fmoc-Cly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Ser(OtBu)-OH, Boc-His(Trt)-OH (Step 1).

The selective deprotection of the Lys(Aloc) group was performed

25 manually and accomplished by treating the resin with a solution of 3 eq
of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h
(Step 2). The resin was then washed with CHCl₃ (6 x 5 mL), 20% HOAc
in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL . The synthesis
was then re-automated for the addition of the two AEEA

30 (aminoethoxyethoxyacetic acid) groups and the 3-maleimidopropionic
acid (Step 3). Resin cleavage and product isolation was performed
using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by
precipitation by dry-ice cold Et₂O (Step 4). The product was purified by
preparative reverse phase HPLC using a Varian (Rainin) preparative

binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH_3CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

Example 34 - Modification of HIV-1 DP 178 at the C-Terminus
10 Preparation of modified HIV-1 DP 178 antifusogenic peptide
Tyr-Thr-Ser-Leu-Ile-His-Ser-Leu-Ile-Glu-Glu-Ser-Gln-Asn-Glu-Glu-Glu-Glu-Leu-Asp-Lys-Trp-Ala-Ser-Leu-Trp-Asn-Trp-Phe-Lys-(Nε-MPA)-NH₂

Using automated peptide synthesis, the following protected amino 15 acids are sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Mtt)-OH, Fmoc-Phe-OH, Fmoc-Trp(Boc)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Trp(Boc)-OH, Fmoc-Leu-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ala-OH, Fmoc-Trp(Boc)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asp(tBu)-OH, Fmoc-Leu-OH, Fmoc-Glu(tBu)-OH, Fmoc-Leu-OH, Fmoc-Leu-OH, Fmoc-Glu(Tbu)-OH, 20 Fmoc-Gln(Trt)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-lle-OH, Fmoc-Leu-OH, Fmoc-Ser(tBu)-OH, Fmoc-His(Trt)-OH, Fmoc-Ile-OH, Fmoc-Leu-OH, Fmoc-25 Ser(tBu)-OH, Fmoc-Thr(tBu)-OH and Boc-Tyr(tBu)-OH. Manual synthesis is employed for the remaining steps: selective removal of the Mtt group and coupling of maleimidopropionic acid (MPA) using HBTU/HOBt/DIEA activation in DMF. The target molecule is removed from the resin; the product is isolated by precipitation and purified by 30 preparative HPLC to afford the desired product as a white solid upon lyophilization.

Example 35 - Modification of HIV-1 DP 107 at the C-Terminus

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Preparation of modified HIV-1 DP 107 antifusogenic peptide Asn-Asn-Leu-Leu-Arg-Ala-IIe-Glu-Ala-Glu-Glu-His-Leu-Leu-Glu-Leu-Thr-Val-Trp-Glu-IIe-Lys-Glu-Leu-Glu-Ala-Arg-IIe-Leu-Ala-Val-Glu-Arg-Tyr-Leu-Lys-Asp-Glu-Lys-(Νε-ΜΡΑ)NΗ₂

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Using automated peptide synthesis, the following protected amino acids are sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Mtt)-OH, Fmoc-Gin(Trt)-OH, Fmoc-Asp(tBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Leu-OH, Fmoc-Ile-10 OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Leu-OH, Fmoc-Gln(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ile-OH, Fmoc-Gln(Trt)-OH, Fmoc-Trp(Boc)-OH, Fmoc-Val-OH, Fmoc-Thr(tBu)-OH, Fmoc-Leu-OH, Fmoc-Gln(Trt)-OH, Fmoc-Leu-OH, Fmoc-Leu-OH, Fmoc-His(Trt)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Ala-OH, 15 Fmoc-Glu(tBu)-OH, Fmoc-Ile-OH, Fmoc-Ala-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Leu-OH, Fmoc-Asn(Trt)-OH, Fmoc-Asn(Trt)-OH. Manual synthesis is employed for the remaining steps: selective removal of the Mtt group and coupling of maleimidopropionic acid (MPA) using HBTU/HOBt/DIEA activation in DMF. The target analog is removed from 20 the resin; the product is isolated by precipitation and purified by preparative HPLC to afford the desired product as a white solid upon lyophilization.

25 2. Modification of the Therapeutic Peptide at the N-Terminus

Example 36 – Modification of RSV Peptide at the ε-Amino Group of the Added N-terminus Lysine Residue Preparation of (Nε-MPA)-Lys-Val-Ile-Thr-Ile-Glu-Leu-Ser-Asn-Ile-Lys-Glu-Asn-Lys-Met-Asn-Gly-Ala-Lys-Val-Lys-Leu-Ile-Lys-Gln-Glu-Leu-Asp-Lys-Tyr-Lys-Asn-Ala-Val

Solid phase peptide synthesis of a modified RSV peptide on a 100 µmole scale is performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin:

Fmoc-Val-OH, Fmoc-Ala-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asp(tBu)-OH, Fmoc-Leu-OH, Fmoc-Glu(tBu)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ile-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Val-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ala-OH, Fmoc-Gly-OH, Fmoc-Asn(Trt)-OH, Fmoc-Met-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Glu(tBu)-OH. Fmoc-Lys(Boc)-OH, Fmoc-Ile-OH, Fmoc-Asn(Trt)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Leu-OH, Fmoc-Glu(tBu)-OH, Fmoc-Ile-OH, Fmoc-Thr(tBu)-OH, Fmoc-Ile-OH, Fmoc-Val-OH, Fmoc-Lys(Aloc)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the 10 sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyluronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is 15 performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh3)4 dissolved in 5 mL of CHCl3:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl3 (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of the 3-20 maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with N,N-dimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et2O (Step 4). The product is purified by preparative reversed 25 phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H2O (A) and 0.045% TFA in CH3CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in 30 >95% purity, as determined by RP-HPLC.

Example 37 – Modification of Neuropeptide Y at the ε-Amino Group of the Added N-terminus Lysine Residue Preparation of (N-εMPA)-Lys-Tyr-Pro-Ser-Lys-Pro-Glu-Asn-Pro-Gly-Glu-Asp-Ala-Pro-Ala-Glu-Asp-Met-Ala-Arg-Tyr-Tyr-Ser-Ala-Leu

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Solid phase peptide synthesis of a modified neuropeptide Y on a 100 µmole scale was performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Leu-OH, Fmoc-Ala-OH, Fmoc-Ser(tBu)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Tvr(tBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ala-OH, Fmoc-Met-OH, Fmoc-Asp(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Ala-OH, Fmoc-Pro-OH, Fmoc-Ala-OH, Fmoc-Asp(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Gly-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Pro-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Lys(Aloc)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl3:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with N,Ndimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian

(Rainin) preparative binary HPLC system: gradient elution of 30-55% B

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 $(0.045\%~TFA~in~H_2O~(A)~and~0.045\%~TFA~in~CH_3CN~(B))$ over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

Example 38 – Modification of Neuropeptide Y at the ε-Amino Group of the Added N-terminus Lysine Residue Preparation of (N-εMPA)-Lys-Tyr-Pro-Ser-Lys-Pro-Asp-Asn-Pro-Gly-Glu-Asp-Ala-Pro-Ala-Glu-Asp-Met-Ala-Arg-Tyr-Tyr-Ser-Ala-Leu

Solid phase peptide synthesis of a modified neuropeptide Y on a 100 µmole scale was performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: 15 Fmoc-Leu-OH, Fmoc-Ala-OH, Fmoc-Ser(tBu)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ala-OH, Fmoc-Met-OH, Fmoc-Asp(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Ala-OH, Fmoc-Pro-OH, Fmoc-Ala-OH, Fmoc-Asp(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Gly-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Asp(tBu)-OH, Fmoc-Pro-OH, 20 Fmoc-Lys(Boc)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Lys(Aloc)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is 25 achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)4 dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The 30 resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 3).

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Between every coupling, the resin is washed 3 times with N,Ndimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et2O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian 5 (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH_3CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

Example 39 – Modification of Dyn A 1-13 at the ε-Amino Group of the Added N-terminus Lysine Residue - Synthesis of (N ϵ -MPA)-Dyn 15 A 1-13-NH₂ $\textbf{(N}\epsilon\text{-MPA)-Lys-Tyr-Gly-Gly-Phe-Leu-Arg-Arg-lle-Arg-Pro-Lys-Leu}$

Solid phase peptide synthesis of a modified Dyn A 1-13 analog on a 100 µmole scale is performed using manual solid-phase synthesis, a 20 Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Pro-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ile-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Phe-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, 25 Fmoc-Tyr(tBu)-OH, Fmoc-Lys(Aloc)-OH. They were dissolved in N,Ndimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 30 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h

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(Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of the 3-maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with *N,N*-dimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC:

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Example 40 - Modification of Dyn A 2-17-NH₂ at the N-terminus Glycine - Synthesis of MPA-AEA₃-Dyn A 2-17-NH₂ (MPA-AEA-AEA)-Gly-Gly-Phe-Leu-Arg-Arg-Ile-Arg-Pro-Lys-Leu-Lys-Trp-Asp-Asn-Glu

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Using automated peptide synthesis, the following protected amino acids and maleimide were sequentially added to Rink Amide MBHA resin: Fmoc-Gln(Trt)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Trp(Boc)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Pro-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Ile-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-AEA-OH, Fmoc-AEA-OH, Fmoc-AEA-OH, Fmoc-AEA-OH, Fmoc-AEA-OH, Fmoc-AEA-OH, Fmoc-AEA-OH, proc-AEA-OH, The target dynorphin analog was then removed from the resin; the product was isolated by precipitation and purified by preparative HPLC to afford the desired product as a pale yellow solid upon lyophilization in a 32% yield. Anal. HPLC indicated product to be >95% pure with $R_t = 33.44$ min. ESI-MS m/z for $C_{109}H_{172}N_{35}O_{29}$ (MH $^+$), calcd 2436.8, found MH $^{3+}$ 813.6.

Example 42 – Modification of Kringle-5 at the ε-Amino Group of the Added N-Terminus Lysine Residue Preparation of (Nε-MPA)-Lys-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂.2TFA

Solid phase peptide synthesis of a modified Kringle-5 analog on a 100 umole scale was performed using manual solid-phase synthesis, a 10 Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Tyr(tBu)-OH, Fmoc-Asp(tBu)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Lys(Aloc)-OH. They were dissolved in N,N-dimethylformamide (DMF) 15 and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (step 1). The selective 20 deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 25 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with N,Ndimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The 30 product is purified by preparative reverse phase HPLC using a Varian

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(Rainin) preparative binary HPLC system: gradient elution of 30-55% B $(0.045\% \text{ TFA in H}_2\text{O} \text{ (A)} \text{ and } 0.045\% \text{ TFA in CH}_3\text{CN (B)})$ over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

Example 43 – Modification of Kringle-5 at the N-Terminus Proline 10 Preparation of (MPA-AEEA)-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lvs(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-15 Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH (step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of Fmoc-AEEA. Deprotection of the resulting Fmoc-AEEA-peptide with piperidine 20% in DMF allow for the subsequent addition of the 3-MPA 20 (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm 25 column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-30 spray ionization.

Example 44 - Modification of Kringle-5 at the N-Terminus Proline Preparation of (MPA)-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-Lys-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-5 Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH (step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation was performed 10 using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization

Example 45 - Modification of Kringle-5 at the N-Terminus Tyrosine Preparation of (MPA-AEEA)-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂.2TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tvr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH (Step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of Fmoc-AEEA. Deprotection of the resulting Fmoc-AEEApeptide with piperidine 20% in DMF allow for the subsequent addition of

the 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 μm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

<u>Example 46</u> - Modification of Kringle-5 at the N-Terminus Tyrosine Preparation of (MPA)-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂.2TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lvs(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, 20 Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH (Step1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% 25 phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian 30 Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard

LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

5 <u>Example 47</u> – Modification of Kringle-5 at the N-Terminus Arginine Preparation of (MPA-AEEA)-Arg-Asn-Pro-Asp-Gly-Asp-Gly-Pro-Trp-Ala-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂.3TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-10 Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Ala-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, 15 Fmoc-Gly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Arg(Pbf)-OH (step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of Fmoc-AEEA. Deprotection of the resulting Fmoc-AEEApeptide with piperidine 20% in DMF allow for the subsequent addition of 20 the 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈. 25 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector 30 and using electro-spray ionization.

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Example 48 – Modification of Kringle-5 at the N-Terminus Arginine Preparation of (MPA)-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-Ala-Tyr-Thr-Thr-Asn-Pro-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂.3TFA

Using automated peptide synthesis, the following protected amino 5 acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tvr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Ala-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, 10 Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Gly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Arg(Pbf)-OH (Step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation was 15 performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 20 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector 25 and using electro-spray ionization.

Example 49 – Modification of Kringle-5 at the N-Terminus Arginine Preparation of (MPA-AEEA)-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Pro-Trp-NH₂.TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Gly-OH, Fmoc-

Asp(OtBu)-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Arg(Pbf)-OH

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(step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of Fmoc-AEEA. Deprotection of the resulting Fmoc-AEEA-peptide with piperidine 20% in DMF allow for the subsequent addition of the 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 μm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

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<u>Example 50</u> – Modification of Kringle-5 at the N-Terminus Arginine Preparation of (MPA)-Arg-Asn-Pro-Asp-Gly-Asp-Val-Gly-Gly-Pro-Trp-NH₂.TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Trp-OH, Fmoc-Pro-OH, Fmoc-Gly-OH, Fmoc-Gly-OH, Fmoc-Cly-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Asp(OtBu)-OH

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254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electrospray ionization.

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Example 51 – Modification of Kringle-5 at the N-Terminus Arginine Preparation of (MPA-AEEA)-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-10 Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH (Step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of Fmoc-AEEA. Deprotection of the resulting Fmoc-AEEA-peptide with piperidine 20% in 15 DMF allow for the subsequent addition of the 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC 20 system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer 25 equipped with a diode array detector and using electro-spray ionization

Example 52 – Modification of Kringle-5 at the N-Terminus Arginine Preparation of (MPA)-Arg-Lys-Leu-Tyr-Asp-Tyr-NH₂.2TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lys(Boc)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Asp(OtBu)-OH, Fmoc-

Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH (Step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 μm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electrospray ionization.

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Example 53 – Modification of Kringle-5 at the N-Terminus Proline Preparation of (MPA-AEEA)-Pro-Arg-Lys-Leu-Tyr-Asp-NH₂.2TFA

Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-20 Lys(Boc)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH (step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of Fmoc-AEEA. Deprotection of the resulting Fmoc-AEEA-peptide with piperidine 20% in 25 DMF allow for the subsequent addition of the 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC 30 system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm.

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The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

5 Example 54 – Modification of Kringle-5 at the N-Terminus Proline Preparation of (MPA)-Pro-Arg-Lys-Leu-Tyr-Asp-NH₂.2TFA

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Using automated peptide synthesis, the following protected amino acids were sequentially added to Rink Amide MBHA resin: Fmoc-Lvs(Boc)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Tyr(tBu)OH, Fmoc-Leu-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH (Step 1). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine (Step 2) followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product was purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization.

3. Modification at an Internal Amino Acid

Example 55 - Synthesis of Lys²⁶(ε-MPA)GLP-1(7-36)-NH₂

Solid phase peptide synthesis of a modified GLP-1 analog on a 5 100 µmole scale was performed manually and on a Symphony Peptide Synthesizer using Fmoc protected Rink amide MBHA resin. The following protected amino acids are sequentially added to the resin: Fmoc-Arg(Pbf)-OH, Fmoc-Gly-OH, Fmoc-Lys(Boc)-OH, Fmoc-Val-OH, Fmoc-Leu-OH, Fmoc-Trp(Boc)-OH, Fmoc-Ala-OH, Fmoc-Ile-OH, Fmoc-10 Phe-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(Aloc)-OH, Fmoc-Ala-OH, Fmoc-Ala-OH, Fmoc-Gln(Trt)-OH, Fmoc-Gly-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Leu-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Val-OH, Fmoc-Asp(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Gly-OH, Fmoc-15 Glu(OtBu)-OH, Fmoc-Ala-OH, Boc-His(Trt)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). Selective deprotection of the Lys(Aloc) group is performed manually and accomplished by treating the resin with a solution of 3eq of Pd(PPh₃)₄ dissolved in 5mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2h (Step 2). The resin is then washed with CHCl₃ (6 x 5mL), 20% HOAc in DCM 25 (6 x 5mL), DCM (6 x 5mL), and DMF (6 x 5mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation is performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative 30 reversed phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column

equipped with a Dynamax C_{18} , 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic diagram below.

Fmoc-Rink Amide MBHA Resin

Boc-HAEGTFTSDVSSYLEGQAA-Lys(Aloc)-EFIAWLVKCR-PS

Boc-HAEGTFTSDVSSYLEGQAA-Lys-EFIAWLVKGR-PS

Step 4 85% TFA/5% TIS/5% thioanisole/5% phenol

Lys²⁶ (E-MPA)GLP-1 (7-36)-NH₂

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D. Preparation of Modified Peptides From Peptides Containing One Free Cysteine

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Preparation of maleimido peptides from therapeutic peptides containing one free Cysteine is exemplified by the synthesis of peptides as described below. The peptide may be modified at the N-terminus, the

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C-terminus, or at an amino acid located between the N-terminus and the C-terminus.

Preparation of maleimido peptides from peptides containing multiple protected functional groups and multiple Cysteine residues all with one free Cysteine residues (*i.e.* all Cysteine residues, except one, are tied up as disulfides). Linking from an internal amino acid in the natural sequence as in Example 5. The free Cysteine residue must be capped or replaced with another amino acid (e.g. Alanine, Methionine, etc.).

Where the peptide contains one cysteine, the cysteine must stay capped after addition of the maleimide. If the cysteine is involved in binding site, assessment has to be made of how much potency is lost is cysteine is capped by a protecting group. If the cysteine can stay capped, then the synthetic path is similar to example (i) above.

Examples of therapeutic peptides that contain one cystein include G_{α} (the alpha subunit of Gtherapeutic peptide binding protein), the 724-739 fragment of rat brain nitric oxide synthase blocking peptide, the alpha subunit 1-32 fragment of human [Tyr0] inhibin, the 254-274 fragment of HIV envelope protein, and P34cdc2 kinase fragment.

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1. Modification at the N-Terminus

Example 56 - Modification of Inhibin Peptide at the Added N-Terminus Lysine

Preparation of (Nε-MPA)-Lys-Tyr-Ser-Thr-Pro-Leu-Met-Ser-Trp-Pro-Trp-Ser-Pro-Ser-Ala-Leu-Arg-Leu-Leu-Gln-Arg-Pro-Pro-Glu-Glu-Pro-Ala-Ala-Ala-His-Ala-Asn-Cys-His-Arg

Solid phase peptide synthesis of a modified inhibin peptide analog on a 100 µmole scale is performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Arg(Pbf)-OH, Fmoc-His(Boc)-OH, Fmoc-Cys(Trt)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Ala-OH, Fmoc-His(Boc)-OH, Fmoc-Ala-

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OH, Fmoc-Ala-OH, Fmoc-Pro-OH, Fmoc-Glu(tBu)-OH, Fmoc-Glu(tBu)-OH, Fmoc-Pro-OH, Fmoc-Pro-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Gln(Trt)-OH, Fmoc-Leu-OH, Fmoc-Leu-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Leu-OH, Fmoc-Ala-OH, Fmoc-Ser(tBu)-OH, Fmoc-Pro-OH, Fmoc-Ser(tBu)-OH, Fmoc-Trp(Boc)-OH, Fmoc-Pro-OH, Fmoc-Trp(Boc)-OH, Fmoc-Ser(tBu)-5 OH, Fmoc-Met-OH, Fmoc-Leu-OH, Fmoc-Pro-OH, Fmoc-Thr(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Lys(Aloc)-OH, They are dissolved in N.N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyluronium hexafluorophosphate (HBTU) and Diisopropylethylamine 10 (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc 15 (18:1:0.5) for 2 h (Step 2). The resin is then washed with $CHCl_3$ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of the 3maleimidopropionic acid (Step 3). Between every coupling, the resin is 20 washed 3 times with N,N-dimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH_3CN 25 (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenylhexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

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<u>Example 57</u> – Modification of RSV Antifusogenic Peptide at the N-Terminus

Preparation of 3-(MPA)-Val-lle-Thr-lle-Glu-Leu-Ser-Asn-lle-Lys-Glu-Asn-Lys-Cys-Asn-Glu-Ala-Lys-Val-Lys-Leu-lle-Lys-Glu-Glu-Leu-Asp-Lys-Tyr-Lys-Asn-Ala-Val

Initially, (Cysteine (Cys) was replaced with Methionine (Met) within the native sequence. Solid phase peptide synthesis of a modified anti RSV analog on a 100 µmole scale was performed on a Symphony Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin, 10 Fmoc protected amino acids, O-benzotriazol-1-yl-N, N, N, N, tetramethyl-uronium hexafluorophosphate (HBTU) in N,Ndimethylformamide (DMF) solution and activation with N-methyl morpholine (NMM), and piperidine deprotection of Fmoc groups (Step 1). The deprotection of the terminal Fmoc group is accomplished using 15 20% piperidine (Step 2) followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation was performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient 20 elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH_3CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenylhexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC. These steps are illustrated by the schematic 25 diagram below.

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Fmoc-Rink Amide MBHA Resin

Fmoc-VITIELSNIKENKMNGAKVKLIKQELDKYKNAV-PS

H, N-VITIELSNIKENKMNGAKVKLIKQELDKYKNAVK-PS

2. Modification at the C-Terminus

Example 58 - Modification of Inhibin Peptide at the Added C-Terminus Lysine Preparation of (Nε-MPA)-Lys-Cys-Asn-Leu-Lys-Glu-Asp-Gly-Ile-Ser-Ala-Ala-Lys-Asp-Val-Lys

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Solid phase peptide synthesis of a modified inhibin peptide analog on a 100 µmole scale is performed using manual solid-phase synthesis, a Symphony Peptide Synthesizer and Fmoc protected Rink Amide MBHA. The following protected amino acids are sequentially added to resin: Fmoc-Lys(Boc)-OH, Fmoc-Val-OH, Fmoc-Asp(tBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Ala-OH, Fmoc-Ala-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ile-OH, Fmoc-Gly-OH, Fmoc-Asp(tBu)-OH, Fmoc-Cys(Trt)-Fmoc-Lys(Boc)-OH, Fmoc-Leu-OH, Fmoc-Asp(Trt)-OH, Fmoc-Cys(Trt)-

OH, Fmoc-Lys(Aloc)-OH, They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-vI-N. N. N'. N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). The selective deprotection of the Lys (Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 10 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 3). Between every coupling, the resin is washed 3 times with N,Ndimethylformamide (DMF) and 3 times with isopropanol. The peptide is cleaved from the resin using 85% TFA/5% TIS/5% thioanisole and 5% 15 phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 µ phenyl-hexyl, 21 mm x 25 20 cm column and UV detector (Varian Dynamax UVD II) at $\lambda\,214$ and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

Example 59 – Modification of RSV Fusogenic Peptide at the CTerminus
Val-Ile-Thr-Ile-Glu-Leu-Ser-Asn-Ile-Lys-Glu-Asn-Lys-Cys-Asn-Gly-Ala-Lys-Val-Lys-Leu- Ile- Lys-Gln-Glu-Leu-Asp-Lys-Tyr-Asn-Ala-Val-(AEEA.MPA)

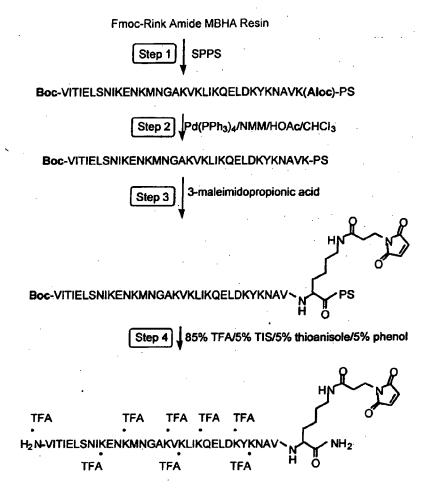
Preparation of maleimido peptides from peptides containing multiple protected functional groups and one cysteine is exemplified by the synthesis of a modified RSV fusogenic peptide. The modified RSV fusogenic peptide was synthesized by linking off the C-terminus by the

addition of a lysine residue to the natural peptide sequence as illustrated by the schematic diagram below. In cases where a cysteine residue is contained within the peptide sequence and is not essentially to the biological activity of the peptide, this residue must be replaced with another amino acid (e.g. alanine, methionine, etc.). In the following synthesis, the cysteine (Cys) was replaced with a methionine (Met) within the native RSV sequence.

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Solid phase peptide synthesis of the maleimido RSV fusogenic peptide on a 100 µmole scale is performed using manual solid-phase synthesis and a Symphony Peptide Synthesizer using Fmoc protected 10 Rink Amide MBHA resin, Fmoc protected amino acids, O-benzotriazol-1yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) in N, Ndimethylformamide (DMF) solution and activation with N-methyl morpholine (NMM), and piperidine deprotection of Fmoc groups (Step 1). The selective deprotection of the Lys(Aloc) group is performed 15 manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of 3-maleimidopropionic acid (Step 3). 20 Resin cleavage and product isolation is performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H_2O (A) and 0.045% TFA in CH_3CN 25 (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenylhexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic 30 diagram below.

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3. Modification at an Internal Amino Acid

5 Example 60 – Modification of $G\alpha$ Peptide at Cys-Asn-Leu-Lys-Glu-Asp-Gly-Ile-Ser-Ala-Ala-Lys-Asp-Val

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Preparation of maleimido peptides from peptides containing multiple protected functional groups and one cysteine is exemplified by the synthesis of a modified $G\alpha$ peptide. The modified $G\alpha$ peptide is synthesized by linking at an internal amino acid as described below.

In cases where a cysteine residue is contained within the peptide sequence and is not essentially to the biological activity of the peptide, this residue must be capped or replaced with another amino acid (e.g. alanine, methionine, etc.). Solid phase peptide synthesis of the modified

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Ga peptide on a 100 µmole scale is performed using manual solidphase synthesis and a Symphony Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin, Fmoc protected amino acids, Obenzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) in N,N-dimethylformamide (DMF) solution and activation with Nmethyl morpholine (NMM), and piperidine deprotection of Fmoc groups (Step 1). The selective deprotection of the Lys(Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq. of Pd(PPh₃)₄ dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2 h (Step 2). The resin is then washed with CHCl₃ (6 x 5 mL), 20% HOAc in DCM (6 x 5 mL), DCM (6 x 5 mL), and DMF (6 x 5 mL). The synthesis is then re-automated for the addition of 3-maleimidopropionic acid (Step 3). Resin cleavage and product isolation is performed using 85% TFA/5% TIS/5% thioanisole and 5% phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reversed phased HPLC using a Varian (Rainin) preparative binary HPLC system: gradient elution of 30-55% B (0.045% TFA in H₂O (A) and 0.045% TFA in CH₃CN (B)) over 180 min at 9.5 mL/min using a Phenomenex Luna 10 μ phenyl-hexyl, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC.

E. Preparation of Modified Peptides From Peptides Containing Two Cysteines In Disulfide Bridge

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Where the peptide contains two cysteines as a disulfide bridge, the peptide is cleaved from the support resin before addition of the maleimide. We need to add a Lys protected with a Mtt group in order to selectively deprotect the lysine in presence of other t-Boc protected lysine. All protecting groups are present except at the carboxy terminus (which stays unprotected due to cleavage from the support resin) and at the two cysteines, which need to be deprotected when peptide is

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cleaved from resin. Mild air oxidation yield the disulfide bridge, and the peptide can be purified at that stage. Solution phase chemistry is then required to activate the C-terminus in presence of the disulfide bridge and add the maleimide (through an amino-alkyl-maleimide) to the C-terminus. The peptide is then fully deprotected. Examples of therapeutic peptides that contain two cysteins as a disulfide bridge include human osteocalcin 1-49, human diabetes associated peptide, the 5-28 fragment of human/canine atrial natriuretic peptide, bovine bactenecin, and human [Tyr0]-cortistatin 29.

Preparation of maleimido peptides from therapeutic peptides containing two Cysteines in a disulfide bridge is exemplified by the synthesis of peptides as described below. The peptide may be modified at the N-terminus, the C-terminus, or at an amino acid located between the N-terminus and the C-terminus.

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1. Modification at the N-Terminus

Example 61 – Modification of TH-1 Peptide at N-Terminus Preparation of (Nε-MPA)NH₂-Lys-Arg-Gly-Asp-Ala-Cys-Glu-Gly-Asp-Ser-Gly-Gly-Pro-Phe-Cys

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Preparation of thiol cyclized maleimido peptides from peptides containing multiple protected functional groups and no free cysteine residues (*i.e.* all cysteine residues are tied up as disulfide bridges). is illustrated by the synthesis of a modified TH-1 peptide.

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Solid phase peptide synthesis of the modified TH-1 peptide on a 100 µmole scale was performed manually and on a Symphony Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin, Fmoc protected amino acids, O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) in N,N-dimethylformamide (DMF) solution and activation with N-methyl morpholine (NMM), and piperidine deprotection of Fmoc groups (Step 1). The removal of the Acm group and resulting oxidation of the two Cys residues to form the cyclized on

resin DAC was accomplished using TI(CF₃CO)₂ (Step 2). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation was performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice 5 cold Et₂O (Step 4). The product was purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm. The 10 product had >95% purity as determined by RP-HPLC mass spectrometry using a Hewlett Packard LCMS-1100 series spectrometer equipped with a diode array detector and using electro-spray ionization, ESI-MS m/z for $C_{66}H_{95}N_{20}O_{26}S_2$ (MH⁺), 1646.8. Found: 1646.7. These steps are illustrated in the schematic diagram below. 15

Fmoc-Rink Amide MBHA Resin

Fmoc-K(Boc)R(Pbf)GD(OtBu)AC(Acm)E(OtBu)GD(OtBu)S(tBu)GGPFC(Acm)-Resin

Step 2

Fmoc-K(Boc)R(Pbf)GD(OtBu)ACE(OtBu)GD(OtBu)S(tBu)GGPFC-Resin

Step 3

1. 20% piperidine
2. 3-maleimidopropionic acid

K(Boc)R(Pbf)GD(OtBu)ACE(OtBu)GD(OtBu)S(tBu)GGPFC-Resin

Step 4

86% TFA/5% TIS/5% H₂O/2% thioanisole/2% phenol

TFA

KRGDACEGDSGGPFC-NH₂

TFA

Example 62 - Synthesis of N-MPA-Ser¹-Somatostatin-28

Solid phase peptide synthesis of the DAC:Somatostatin-28 analog on a 100 µmole scale is performed manually and on a Symphony 5 Peptide Synthesizer using Fmoc protected Rink amide MBHA resin. The following protected amino acids are sequentially added to the resin: Fmoc-Cys(Acm)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Trp(Boc)-OH, Fmoc-Phe-OH, Fmoc-Phe-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, 10 Fmoc-Cys(Acm)-OH, Fmoc-Gly-OH, Fmoc-Ala-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Ala-OH, Fmoc-Met-OH, Fmoc-Ala-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Ala-OH, Fmoc-Ser(tBu)-OH. They are dissolved in N,N-dimethylformamide 15 (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). Removal of the Acm 20 groups and resulting oxidation of the two Cys residues to form the disulfide bridge is accomplished using iodine (Step 2). Deprotection of the terminal Fmoc group is accomplished using 20% piperidine followed by the coupling of 3-MPA (Step 3). Resin cleavage and product isolation is performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% 25 phenol, followed by precipitation by dry-ice cold Et₂O (Step 4). The product is purified by preparative reversed phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 μm guard module, 21 mm x 25 cm column and UV detector (Varian 30 Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in

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>95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic diagram below.

Fmoc-Rink Amide MBHA Resin

Step 1 SPPS

Fmoc-SANSNPAMAPRERKAGC(Acm)KNFFWKTFTSC(Acm)-Resin

Step 2 12

Fmoc-SANSNPAMAPRERKAGCKNFFWKTFTSC-Resin

Step 3 2 2. 3-maleimidopropionic acid

Step 4 85% TFA/5% TIS/5% thioanisole/5% pheno

N-MPA-Ser 1-Somatostatin-28

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2. Modification at the C-Terminus

Example 63 - Synthesis of Somatostatin-28-EDA-MPA

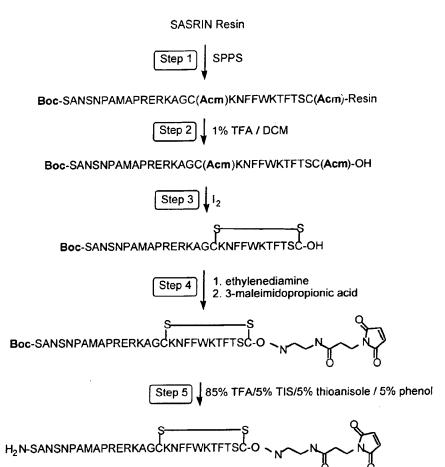
Solid phase peptide synthesis of the DAC:Somatostatin-28

analog on a 100 µmole scale is performed manually and on a Symphony Peptide Synthesizer using SASRIN (super acid sensitive resin). The following protected amino acids are sequentially added to the resin:

Fmoc-Cys(Acm)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Cys(Boc)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Phe-OH, Fmoc-Phe-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Cys(Acm)-OH, Fmoc-Gly-OH, Fmoc-Ala-OH, Fmoc-Lys(Boc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Fmoc-Cys(Pbf)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Cys(Pbf)-OH, F

Pro-OH, Fmoc-Ala-OH, Fmoc-Met-OH, Fmoc-Ala-OH, Fmoc-Pro-OH. Fmoc-Asn(Trt)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Ala-OH. Boc-Ser(tBu)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-vl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). The fully protected peptide is cleaved from the resin by treatment with 1% TFA / DCM (Step 2). Removal of the Acm groups and resulting oxidation of the two Cys 10 residues to form the disulfide bridge is accomplished using iodine (Step 3). Ethylenediamine and 3-maleimidopropionic acid are then sequentially added to the free C-terminus (Step 4). The protecting groups are then cleaved and the product isolated using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O 15 (Step 5). The product is purified by preparative reversed phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the 20 desired DAC in >95% purity, as determined by RP-HPLC. These steps

are illustrated in the schematic diagram below.



Somatostatin-28-EDA-MPA

3. Modification at an Internal Amino Acid

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Example 64 – Synthesis of Lys¹⁴(ε-MPA)-Somatostatin-28

Solid phase peptide synthesis of the DAC:Somatostatin-28 analog on a 100 µmole scale is performed manually and on a Symphony Peptide Synthesizer using Fmoc protected Rink amide MBHA resin. The following protected amino acids are sequentially added to the resin: Fmoc-Cys(Acm)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Phe-OH, Fmoc-Thr(tBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Trp(Boc)-OH,

Fmoc-Phe-OH, Fmoc-Phe-OH, Fmoc-Asn(Trt)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Cys(Acm)-OH, Fmoc-Gly-OH, Fmoc-Ala-OH, Fmoc-Lys(Aloc)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Arg(Pbf)-OH, Fmoc-Pro-OH, Fmoc-Ala-OH, Fmoc-Met-OH, Fmoc-Ala-OH, Fmoc-Pro-OH, Fmoc-Asn(Trt)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Asn(Trt)-OH, Fmoc-Ala-OH, Fmoc-Ser(tBu)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,N-10 dimethylformamide (DMF) for 20 minutes (Step 1). Removal of the Acm groups and resulting oxidation of the two Cys residues to form the disulfide bridge is accomplished using iodine (Step 2). Selective deprotection of the Lys(Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh₃)₄ 15 dissolved in 5mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2h (Step 3). The resin is then washed with CHCl₃ (6 x 5mL), 20% HOAc in DCM (6 x 5mL), DCM (6 x 5mL), and DMF (6 x 5mL). The synthesis is then reautomated for the addition of the 3-maleimidopropionic acid (Step 4). Resin cleavage and product isolation is performed using 86% TFA/5% 20 TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 5). The product is purified by preparative reversed phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm 25 column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC. These steps are illustrated in the following schematic diagram.

Fmoc-Rink Amide MBHA Resin

Boc-SANSNPAMAPRER-Lys(Aloc)AGC(Acm)KNFFWKTFTSC(Acm)-Resin

Boc-SANSNPAMAPRER-Lys(Aloc)AGCKNFFWKTFTSC-Resin

Boc-SANSNPAMAPRER-Lys-AGCKNFFWKTFTSC-Resin

Step 5 \$\int 85\% TFA/5\% TIS/5\% thioanisole/5\% phenol

Lys¹⁴(E-MPA)-Somatostatin-28

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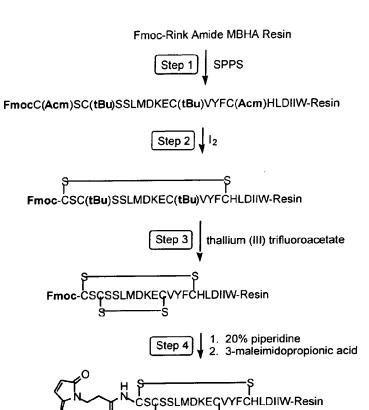
F. Preparation of Modified Peptides From Peptides Containing Multiple Cysteines

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1. Modification at the N-Terminus

Example 65- Synthesis of N-MPA-Cys¹-Endothelin-1 (1-21)-OH

Solid phase peptide synthesis of a modified Endothelin-1 analog on a 100 µmole scale is performed manually and on a Symphony Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin. The 10 following protected amino acids are sequentially added to the resin: Fmoc-Trp(Boc)-OH, Fmoc-Ile-OH, Fmoc-Ile-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Leu-OH, Fmoc-His(Trt)-OH, Fmoc-Cys(Acm)-OH, Fmoc-Phe-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Val-OH, Fmoc-Cys(tBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Met-15 OH, Fmoc-Leu-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Cys(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Cys(Acm)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyluronium hexafluorophosphate (HBTU) and Diisopropylethylamine 20 (DIEA). Removal of the Fmoc protecting groups is achieved using a solution of 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The removal of the Acm groups and resulting oxidation of the first two Cys residues to form the first disulfide bridge on resin is accomplished using iodine (Step 2). The removal of the tBu 25 groups and resulting oxidation of the other two Cys residues to form the second disulfide bridge on resin is accomplished using thallium (III) trifluoroacetate (Step 3). The deprotection of the terminal Fmoc group is accomplished using 20% piperidine followed by the coupling of 3-MPA (Step 4). Resin cleavage and product isolation is performed using 86% 30 TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 5). The product is purified by



N-MPA-Cys1-Endothelin-1 (1-21)-OH

5 preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 μm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 μm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and

254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic diagram above.

Modification at the C-Terminus 2.

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Example 66 - Synthesis of Endothelin-1 (1-21)Lys²²-(Nε-MPA)-OH 5

Solid phase peptide synthesis of a modified Endothelin-1 analog on a 100 umole scale is performed manually and on a Symphony Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin. The following protected amino acids are sequentially added to the resin: Fmoc-Lys(Aloc)-OH, Fmoc-Trp(Boc)-OH, Fmoc-Ile-OH, Fmoc-Ile-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Leu-OH, Fmoc-His(Trt)-OH, Fmoc-Cys(Acm)-OH, Fmoc-Phe-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Val-OH, Fmoc-Cys(tBu)-OH, Fmoc-Glu(OtBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Met-OH, Fmoc-Leu-OH, Fmoc-Ser(tBu)-OH, Fmoc-Ser(tBu)-OH, Fmoc-Cys(tBu)-OH, Fmoc-Ser(tBu)-OH, Boc-Cys(Acm)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N, N-tetramethyl-uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,Ndimethylformamide (DMF) for 20 minutes (Step 1). The removal of the Acm groups and resulting oxidation of the first two Cys residues to form the first disulfide bridge on resin is accomplished using iodine (Step 2). The removal of the tBu groups and resulting oxidation of the other two Cys residues to form the second disulfide bridge on resin is accomplished using thallium (III) trifluoroacetate (Step 3). Selective deprotection of the Lys(Aloc) group is performed manually and accomplished by treating the resin with a solution of 3 eq of Pd(PPh3)4 dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2h (Step 4). The resin is then washed with CHCl3 (6 x 5mL), 20% HOAc in DCM (6 x 5mL), DCM (6 x 5mL), and DMF (6 x 5mL). The synthesis is then re-

automated for the addition of the 3-maleimidopropionic acid (Step 5).

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Resin cleavage and product isolation is performed using 86% TFA/5% TIS/5% $H_2O/2\%$ thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et_2O (Step 5). The product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C_{18} , 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C_{18} , 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic diagram below.

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Fmoc-Rink Amide MBHA Resin

Boc-C(Acm) SC(tBu) SSLMDKEC(tBu) VYFC(Acm) HLDIIW-Lys(Aloc)-Resin

Endothelin-1 (1-21) Lys²²(E-MPA)-OH

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3. Modification at an Internal Amino Acid

Example 67- Synthesis of Lys⁴(Nε-MPA)Sarafotoxin B(1-21)-OH

Solid phase peptide synthesis of a modified Sarafotoxin-B analog on a 100 µmole scale is performed manually and on a Symphony 5 Peptide Synthesizer using Fmoc protected Rink Amide MBHA resin. The following protected amino acids are sequentially added to the resin: Fmoc-Trp(Boc)-OH, Fmoc-Ile-OH, Fmoc-Val-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Gln(Trt)-OH, Fmoc-His(Trt)-OH, Fmoc-Cys(Acm)-OH, Fmoc-Phe-OH, Fmoc-Tyr(tBu)-OH, Fmoc-Leu-OH, Fmoc-Cys(tBu)-OH, Fmoc-10 Glu(OtBu)-OH, Fmoc-Lys(Boc)-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Thr(tBu)-OH, Fmoc-Met-OH, Fmoc-Asp(OtBu)-OH, Fmoc-Lys(Aloc)-OH, Fmoc-Cys(tBu)-OH, Fmoc-Ser(tBu)-OH, Boc-Cys(Acm)-OH. They are dissolved in N,N-dimethylformamide (DMF) and, according to the sequence, activated using O-benzotriazol-1-yl-N, N, N', N'-tetramethyl-15 uronium hexafluorophosphate (HBTU) and Diisopropylethylamine (DIEA). Removal of the Fmoc protecting group is achieved using a solution of 20% (V/V) piperidine in N,N-dimethylformamide (DMF) for 20 minutes (Step 1). The removal of the Acm groups and resulting oxidation of the first two Cys residues to form the first disulfide bridge on 20 resin is accomplished using iodine (Step 2). The removal of the tBu groups and resulting oxidation of the other two Cys residues to form the second disulfide bridge on resin is accomplished using thallium (III) trifluoroacetate (Step 3). Selective deprotection of the Lys(Aloc) group is performed manually and accomplished by treating the resin with a 25 solution of 3 eq of Pd(PPh3)4 dissolved in 5 mL of CHCl₃:NMM:HOAc (18:1:0.5) for 2h (Step 4). The resin is then washed with CHCl3 (6 x 5mL), 20% HOAc in DCM (6 x 5mL), DCM (6 x 5mL), and DMF (6 x 5mL). The synthesis is then re-automated for the addition of the 3maleimidopropionic acid (Step 5). Resin cleavage and product isolation 30 is performed using 86% TFA/5% TIS/5% H₂O/2% thioanisole and 2% phenol, followed by precipitation by dry-ice cold Et₂O (Step 5). The

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product is purified by preparative reverse phase HPLC using a Varian (Rainin) preparative binary HPLC system using a Dynamax C₁₈, 60Å, 8 µm, 21 mm x 25 cm column equipped with a Dynamax C₁₈, 60Å, 8 µm guard module, 21 mm x 25 cm column and UV detector (Varian

Dynamax UVD II) at λ 214 and 254 nm to afford the desired DAC in >95% purity, as determined by RP-HPLC. These steps are illustrated in the schematic diagram below.

Fmoc-Rink Amide MBHA Resin

 $\textbf{Boc-C(Acm)} SC(\textbf{tBu)-Lys}(\textbf{Aloc}) - DMTDKEC(\textbf{tBu}) LYFC(\textbf{Acm}) \\ HQDVIW-Resin$

Step 4 Pd(PPh₃)₄/NMM/HOAc/CHCh

Step 5 3-maleimidopropionic acid

Step 6 \$\int 85\% TFA/5\% TIS/5\% thioanisole/5\% phenol

Lys4(E-MPA)Sarafotoxin B (1-21)-OH

F. Peptide Stability Assays

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Example 68 - Peptide Stability Assay of K5

A peptide stability assay was performed. (MPA)-Pro-Arg-Lys-Leu-Tyr-Asp-Lys-NH₂. 2TFA was synthesized as described above and was identified MPA-K5. The non-modified counterpart peptide Pro-Arg-Lys-Leu-Tyr-Asp-Lys was also synthesized as described in Example 20 without the addition of 3-MPA and identified as K5.

K5 (MW1260.18, 918.12 freebase) was prepared as a 100 mM stock solution in water. MPA-K5 (MW = 1411.17, 1069.11 freebase) was prepared as a 100 mM stock solution in water. Human Serum Albumin (HSA) was obtained as a 25% solution (ca 250 mg/ml, 3.75 mM) as Albutein® available from Alpha Therapeutic. Human plasma was obtained from Golden West Biologicals.

(1) Stability of K5 in Human Plasma

K5 was prepared as a 1 μ M solution and dissolved in 25% human serum albumin. The mixture was then incubated at 37°C in the presence of human plasma to final concentration of 160 mM K5. Aliquots of 100 μ l were withdrawn from the plasma at 0, 4 hours and 24 hours. The 100 μ l aliquots were mixed with 100 μ l of blocking solution (5 vol. 5%ZnSO₄/3 vol. Acetonitrile/2 vol. Methanol) in order to precipitate all proteins. The sample was centrifuged for 5 min at 10,000 g and the supernatant containing the peptide was recovered and filtered through a 0.22 μ m filter. The presence of free intact K5 peptide was assayed by the HPLC/MS. The results are presented below. The HPLC parameters for detection of K5 peptide in serum were as follows.

The HPLC method was as follows: A Vydac C18 250 X 4.6 mm, 5 µ particle size column was utilized. The column temperature was 30°C

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with a flow rate of 0.5 ml/min. Mobile Phase A was 0.1% TFA/water. Mobile Phase B was 0.1% TFA/acetonitrite. The injection volume was 10µl.

The gradient was as follows:

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	Time(Minutes)	<u>%A</u>	<u>%B</u>
	0	95	5
	20	70	30
	25	10	90
10	30	10	90
	35	95	5
	45	95	5

The proteins were detected at 214, 254 and 334 nm. For mass spectral analysis, the ionization mode was API-electrospray (positive mode) at an M/Z range of 300 to 2000. The gain was 3.0, fragmentor 120v, threshold 20, stepsize 0.1. The gas temp was 350°C and the drying gas volume was 10.0 l/min. The Neb pressure was 24 psi and the Vcap was 3500V. The HPLC method was as follows: A Vydac C18 250 X 4.6 mm, 5 µ particle size column was utilized. The column temperature was 30°C with a flow rate of 0.5 ml/min. Mobile Phase A was 0.1% TFA/water. Mobile Phase B was 0.1% TFA/acetonitrite. The injection volume was 10µl.

The gradient was as follows:

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	Time(Minutes)	<u>%A</u>	<u>%B</u>
	0	95 [*]	5
	20	70	30
	25	10	90
30	30	10	90
	35	95	5
	45	95	5

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The proteins were detected at 214, 254 and 334 nm. For mass spectral analysis, the ionization mode was API-electrospray (positive mode) at an M/Z range of 300 to 2000. The gain was 3.0, fragmentor 120v, threshold 20, stepsize 0.1. The gas temp was 350°C and the drying gas volume was 10.0 l/min. The Neb pressure was 24 psi and the Vcap was 3500V.

	<u>Time</u>	%K5 peptide in Plasma
10	0 hrs.	100%
	4 hrs	9%
	24 hrs	0%

The results demonstrate that unmodified K5 peptide is unstable in plasma likely as a result of protease activity.

(2) Stability of MPA-K5-HSA Conjugate in Plasma

MPA-K5 (modified K5 peptide) was incubated with 25% HSA for 2 hours at room temperature. The MPA-K5-HSA conjugate was then incubated at 37° in the presence of human plasma at a final concentration of 160 μ m. After the specific incubation period (0, 4 and 24 hours) an aliquot of 100 μ l was withdrawn and filtered through a 0.22 μ m filter. The presence of intact conjugate was assayed by HPLC-MS.

The column was an Aquapore RP-300, 250 x 4.6 mm, 7µ particle size. The column temperature was 50° C. The mobile phase A was 0.1% TFA/water. The mobile phase B was 0.1% TFA/acetonitrile. The injection volume was 1 µl. The gradient was as follows:

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	Time (minutes)	<u>%A</u>	<u>%B</u>	Flow(ml/min)
	0	66	34	0.700
	1	66	34	0.700
5	25	58.8	41.2	0.700
	30	50	50	0.70
	35	5	95	1.00
	41	5	95	1.00
	45	66	34	1.00
10	46	66	34	0.70

The peptide was detected at 214 mm for quantification. For mass spectral analysis of the peptide, the ionization mode was API-electrospray at 1280 to 1500 m/z range, gain 1.0, fragmentor 125V, threshold 100, stepsize 0.40. The gas temperature was 350°C the drying gas was 13.0 l/min. The pressure was 60psi and the Vcap was 6000V. The results are presented below.

Approximately 33% of circulating albumin in the bloodstream is mercaptalbumin (SH-albumin) which is not blocked by endogenous sulfhydryl compounds such as cysteine or glutathione and is therefore available for reaction with maleimido groups. The remaining 66% of the circulating albumin is capped or blocked by sulfhydryl compounds. The HPLC MS assay permits the identification of capped-HSA, SH-albumin and K5-MPA-albumin. The MPA covalently bonds to the free thiol on the albumin. The stability of the three forms of albumin in plasma is presented below.

	<u>Time</u>	%capped HSA	% SH-Albumin	%K5-MPA-HSA
	0 hrs.	61.3	16.6	22.1
30	4 hrs.	64.6	16.05	19.35
	24 hrs.	63	16.8	20.2

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The percentage of K5-MPA-HSA remained relatively constant throughout the 24 hour plasma assay in contrast to unmodified K5 which decreased to 9% of the original amount of K5 in only 4 hours in plasma. The results demonstrate that in contrast to K5 which is quite unstable in plasma, K5-MPA-HSA is quite stable from peptidase activity in plasma.

Example 70 - Peptide Stability Assay of Dynorphin

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In order to determine the stability of peptide conjugates in the presence of serum peptidases the serum stability of Dyn A-(1-13)-OH, Dyn A-(1-13)-NH₂ and Dyn A 1-13(MPA)-NH₂ were compared. Dyn A-(1-13)-OH, Dyn A-(1-13)-NH₂ and Dyn A 1-13(MPA)-NH₂ were synthesized as described above. The Dynorphin peptides were mixed with human heparinized plasma to a final concentration of 4 mg/mL. After the required incubation time at 37 °C, 0, 20, 20, 60, 120, 180, 360 and 480 minutes) a 100 μ L-aliquot was added to 100 μ L of blocking solution (5 vol. of a 5% aqueous ZnSO₄ solution, 3 vol. of acetonitrile, 2 vol. of methanol) that precipitates all proteins. After centrifugation (10,000 g for 2.5 min), clear supernatant was recovered, filtered through a 0.45 μ m filter and stored on ice until LC/MS analysis.

The samples were analyzed using an LC at 214 nm to detect the presence of the different compounds and MS to determine the identity of the detected compound. The integrated area % for each peak from the LC chromatogram was then plotted against time and the relative stabilities determined in human plasma.

The stability data for Dyn A-(1-13)-OH and Dyn A-(1-13)-NH₂ were consistent with that reported in literature: the proteolytic breakdown of the dynorphin peptides is quite rapid. Dyn A-(1-13)-OH had a half life of about 10 minutes. Dyn A-(1-13)-NH₂ had a half life of about 30 minutes. In contrast Dyn A 1-13(MPA)-NH₂ exhibited striking stabilization in the presence of serum peptidase activity. Unmodified dynorphin peptides are degraded within 60 minutes. In contrast, modified dynorphin

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peptides (Dyn A $1-13(MPA)-NH_2$) are stable from serum peptidase activity for up to 480 minutes.

The stability determination of the dynorphin conjugate is determined by ELISA. In order to determine if the observed signal is due to a dynorphin conjugate and what the conjugate is, LC mass spectrometrytral analysis of the reaction mixture after 8 h was performed. The use of mass spectrometry permits a determination of the molecular weight of the conjugate and allows the determination whether there are any truncated forms of the dynorphin conjugate. Mass spectrometry of human plasma shows the two forms of albumin, the free thiol at 66436 Da and the oxidized form at 66557 Da. Also, mass spectrometry can distinguish between a Dyn 2-13 truncated conjugate (68046 Da) and the intact Dyn 1-13 conjugate, (68207 Da) in an equal mixture.

Mass spectrometry analysis of dynorphin samples taken from the serum after 480 minutes of exposure to the serum peptidases identifies only the presence of the intact conjugate (68192 Da) and not the breakdown products thereby demonstrating the stability of the dynorphin conjugate from serum peptidase activity.

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TABLE 1 NATURAL AMINO ACIDS AND THEIR ABBREVIATIONS			
Alanine	Ala	Α	Fmoc-Ala-OH
Arginine	Arg	R	Fmoc-Arg(Pbf)-OH
Asparagine	Asn	N	Fmoc-Asn(Trt)-OH
Aspartic acid	Asp	D	Asp(tBu)-OH
Cysteine	Cys	С	Fmoc-Cys(Trt)
Glutamic acid	Glu	Е	Fmoc-Glu(tBu)-OH
Glutamine	Gln	Q	Fmoc-Gln(Trt)-OH
Glycine	Gly	G	Fmoc-Gly-OH
Histidine	His	Н	Fmoc-His(Trt)-OH
Isoleucine	lle	ı	Fmoc-Ile-OH
Leucine	Leu	L	Fmoc-Leu-OH
Lysine	Lys	K	Fmoc-Lys(Mtt)-OH
Methionine	Met	M	Fmoc-Met-OH
Phenylalanine	Phe	F	Fmoc-Phe-OH
Proline	Pro	Р	Fmoc-Pro-OH
Serine	Ser	S	Fmoc-Ser(tBu)-OH
Threonine	Thr	Т	Fmoc-Thr(tBu)-OH
Tryptophan	Тгр	W	Fmoc-Trp(Boc)-OH
Tyrosine	Tyr	Υ	Boc-Tyr(tBu)-OH
Valine	Val	V	Fmoc-Val-OH

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We claim:

1. A modified therapeutic peptide capable of forming a peptidase stablilized therapeutic peptide composed of between 3 and 50 amino acids, said peptide having a carboxy terminal amino acid, an amino terminal amino acid, a therapeutically active region of amino acids and a less therapeutically active region of amino acids, said peptide comprising:

a reactive group which reacts with amino groups, hydroxyl
groups, or thiol groups on blood components to form a stable covalent
bond thereby forming the peptidase stabilized therapeutic peptide
wherein the reactive group is selected from the group consisting of
succinimidyl and maleimido groups and wherein the reactive group is
attached to an amino acid positioned in said less therapeutically active
region of amino acids.

2. The peptide of claim 1 wherein said therapeutically active region of amino acids includes said carboxy terminal amino acid and said reactive group is attached to said amino terminal amino acid.

- 3. The peptide of claim 1 wherein said therapeutically active region of amino acids includes said amino terminal amino acid and said reactive group is attached to said carboxy terminal amino acid.
- 25 4. The peptide of claim 1 wherein said therapeutically active region of amino acids includes said carboxy terminal amino acid and said reactive group is attached to an amino acid positioned between said
 - amino terminal amino acid and said carboxy terminal amino acid.
- 30 5. The peptide of claim 1 wherein said therapeutically active region of amino acids includes said amino terminal amino acid and said

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reactive group is attached to an amino acid positioned between said amino terminal amino acid and said carboxy terminal amino acid.

- 6. A method of synthesizing the modified therapeutic peptide of claim 1, comprising:
- a) if said therapeutic peptide does not contain a cysteine, then synthesizing said peptide from said carboxy terminal amino acid and adding said reactive group to said carboxy terminal amino acid, or adding a terminal lysine to said carboxy terminal amino acid and adding said reactive group to said terminal lysine;
- b) if said therapeutic peptide contains only one cysteine, then reacting said cysteine with a protective group prior to addition of said reactive group to an amino acid in said less therapeutically active region of said peptide;
- c) if said therapeutic peptide containe two cysteines as a disulfide bridge, then oxidizing said two cysteines and adding said reactive group to said amino terminal amino acid, or to said carboxy terminal amino acid, or to an amino acid positioned between said carboxy terminal amino acid and said amino terminal amino acid of said therapeutic peptide; and
- d) if said therapeutic peptide contains more than two cysteines as disulfide bridges, then sequentially oxidizing said cysteines in said disulfide bridges and purifying said peptide prior to the addition of said reactive groups to said carboxy terminal amino acid.
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- 7. A method for protecting a therapeutic peptide from peptidase activity in vivo, said peptide being composed of between 3 and 50 amino acids and having a carboxy terminus and an amino terminus and a carboxy terminal amino acid amino acid and an amino terminal amino acid, comprising:
- (a) modifying said peptide by attaching a reactive group to the carboxy terminal amino acid, to the amino terminal amino acid, or to an

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amino acid located between the amino terminal amino acid and the carboxy terminal amino acid, such that said modified peptide is capable of forming a covalent bond in vivo with a reactive functionality on a blood component;

- 5 (b) forming a covalent bond between said reactive group and a reactive functionality on a blood component to form a peptide-blood component conjugate, thereby protecting said peptide from peptidase activity; and
- (c) analyzing the stability of said peptide-blood component
 10 conjugate to assess the protection of said peptide from peptidase activity.
 - 8. A method according to claim 7, further comprising the step of administering said modified peptide in vivo before step (b), such that the peptide-blood component conjugate is formed in vivo.
 - 10. A method according to claim 7, wherein step (b) occurs ex vivo.
- 11. A method according to claim 7 wherein step (c) is performed in 20 vivo.
 - 12. A method according to claim 7, wherein said reactive group is a maleimido group.
 - 13. A method according to claim 7, wherein said reactive group is attached to said peptide via a linking group.
- 30 14. A method according to claim 7, wherein said blood component is albumin.

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15. A method according to claim 7, wherein one or more of said amino acids is synthetic.

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- 16. A method for protecting a therapeutic peptide from peptidase activity in vivo, said peptide being composed of between 3 and 50 amino acids and having a therapeutically active region of amino acids and a less therapeutically active region of amino acids, comprising:
- (a) determining said therapeutically active region of amino acids;
- 10 (b) modifying said peptide at an amino acid included in said less therapeutically active region of amino acids by attaching a reactive group to said amino acid to form a modified peptide, such that said modified peptide has therapeutic activity and is capable of forming a covalent bond in vivo with a reactive functionality on a blood component;
- 15 (c) forming a covalent bond between said reactive entity and a reactive functionality on a blood component to form a peptide-blood component conjugate, thereby protecting said peptide from peptidase activity; and
- (d) analyzing the stability of said peptide-blood component
 20 conjugate to assess the protection of said peptide from peptidase activity.
 - 17. A method according to claim 16, further comprising the step of administering said modified peptide in vivo before step (c), such that the peptide-blood component conjugate is formed in vivo.
 - 18. A method according to claim 18, wherein step (c) occurs ex vivo.
- 19. A method according to claim 19, wherein step (d) is performed in vivo.

- 20. A method according to claim 18, wherein step (c) is performed in vivo.
- 21. A method according to claim 20, wherein step (d) is performed ex 5 vivo.
 - 22. A method according to claim 16, wherein said peptide has a carboxy terminus, an amino terminus, a carboxy terminal amino acid and an amino terminal amino acid, and wherein step (b) further comprises:
- (a) if said less therapeutically active portion is located at the carboxy terminus of said peptide, then modifying said peptide at the carboxy terminal amino acid of said peptide;
 - (b) if said less active portion is located at the amino terminus of said peptide, then modifying said peptide at the amino terminal amino acid of said peptide; and
 - (c) if said less active portion is located at neither the amino terminus nor the carboxy terminus of said peptide, then modifying said peptide at an amino acid located between the carboxy terminus and the amino terminus.

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- 23. A method according to claim 16, wherein said reactive group is a maleimido group.
- 24. A method according to claim 16, wherein said reactive entity is25 attached to said peptide via a linking group.
 - 25. A method according to claim 16, wherein said blood component is albumin.
- 30 26. A method according to claim 16, wherein one or more of said amino acids is synthetic.

SEQUENCE LISTING

<110> Conjuchem, Inc.
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Milner, Peter
Holmes, Darren
Thibaudeau, Karen

<120> PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD COMPONENTS

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Cys Tyr Phe Gln Asn Cys Pro Arg Gly
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Tyr Phe Val Asn Cys Pro Arg Gly
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 Tyr Phe Gln Asn Cys Pro Arg Gly
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 Glu Asn Cys Cys Pro Arg
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Phe Phe Ile Asn Cys Pro Arg Ala
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Cys Tyr Phe Gln Asn Cys Pro Lys Gly
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Tyr Phe Val Asn Cys Pro Gly
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Cys Phe Ile Gln Asn Cys Pro Gly
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 Cys Tyr Ile Gln Asn Cys Pro Arg Gly
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 Tyr Ile Gln Asn Cys Pro Gly
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Ala Trp Pro Ser Pro Ser Val Glu Gly
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<211> 28
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Phe His Leu Leu Arg Glu Val Leu Glu Ala Arg Ala Glu Gln Leu Ala
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Gln Glu Ala His Lys Asn Arg Lys Leu Glu Ile Ile
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Ala Glu Gln Ala Ala Leu Asn Arg Leu Leu Leu Glu Glu Ala
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<211> 41
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Glu Val Leu Glu Met Thr Lys Ala Asp Gln Leu Ala Gln Gln Ala His
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Asn Asn Arg Lys Leu Leu Asp Ile Ala
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<210> 76
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                                   10
Glu Val Leu Glu Met Ala Arg Ala Glu Gln Leu Ala Gln Gln Ala His
            20
                            25
Ser Asn Arg Lys Leu Met Glu Ile Ile
        35
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<210> 77
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<212> PRT
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<223> Description of Artificial Sequence: Synthetic Peptide <400> 77 Ser Gln Glu Pro Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu Arg 10 5 Glu Val Leu Glu Met Thr Lys Ala Asp Gln Leu Ala Gln Gln Ala His 25 Ser Asn Arg Lys Leu Leu Asp Ile Ala 35 <210> 78 <211> 41 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 78 Ser Glu Glu Pro Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu Arg 5 10 Glu Val Leu Glu Met Ala Arg Ala Glu Gln Leu Ala Gln Gln Ala His 25 Ser Asn Arg Lys Leu Met Glu Asn Phe 35 . <210> 79 <211> 21 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 79 Ser Glu Glu Pro Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu Arg 10 15

Glu Val Leu Glu Cys

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<210> 80
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Glu Gln Glu Ala Glu Gln Ala Ala Leu Asn Arg Leu Leu Glu Glu
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Ala
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Phe His Leu Leu Arg Glu Val Leu Glu Ala Arg Ala Glu Gln Leu Ala
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Gln Gln Ala His Ser Asn Arg Lys Leu Glu Ile Ile
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Phe His Leu Leu Arg Glu Val Leu Glu Ala Arg Ala Glu Gln Leu Ala
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Gln Glu Ala His Lys Asn Arg Lys Leu Glu Ile Ile
                               25
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Ser Glu Glu Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu Arg Glu
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Val Leu Glu Met Ala Arg Ala Glu Gln Leu Ala Gln Gln Ala His Ser
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Asn Arg Lys Leu Met Glu Ile Ile
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Ala Gly Ser Glu
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Ser Gln Glu Pro Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu Arg
                 5
                                    10
Glu Val Leu Glu Met Thr Lys Ala Asp Gln Leu Ala Gln Gln Ala His
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             20
                                 25
Ser Asn Arg Lys Leu Leu Asp Ile Ala
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Peptide

<400> 88

Ser Gln Glu Pro Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu Arg

1 5 10 15

Glu Val Leu Glu Thr Lys Ala Asp Gln Leu Ala Gln Gln Ala Tyr Ser 20 25 30

Asn Arg Lys Leu Leu Asp Ile Ala 35 40

<210> 89

<211> 27

<212> PRT

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<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 89

Ser Leu Asp Ser Pro Ala Ala Leu Ala Glu Arg Gly Ala Arg Asn Ala 1 5 10 15

Leu Gly Gly His Gln Glu Ala Pro Glu Arg Glu 20 25

<210> 90

<211> 40

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 90

Glu Gly Pro Pro Ile Ser Ile Asp Leu Ser Leu Glu Leu Leu Arg Lys
1 5 10 15

Met Ile Glu Ile Glu Lys Gln Glu Lys Glu Lys Gln Gln Ala Ala Asn 20 25 30

Asn Arg Leu Leu Leu Asp Thr Ile

35 40

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<211> 42
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Tyr Ser Glu Glu Pro Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu
                          10
Arg Glu Val Leu Glu Met Ala Arg Ala Glu Gln Leu Ala Gln Gln Ala
                   25
His Ser Asn Arg Lys Leu Met Glu Ile Ile
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<211> 42
<212> PRT
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Tyr Ser Gln Glu Pro Pro Ile Ser Leu Asp Leu Thr Phe His Leu Leu
                                  10
Arg Glu Val Leu Glu Met Thr Lys Ala Asp Gln Leu Ala Gln Gln Ala
                       25
                                                  30
His Ser Asn Arg Lys Leu Leu Asp Ile Ala
        35
<210> 93
<211> 9
<212> PRT
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<210> 91

31

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<223> Description of Artificial Sequence: Synthetic
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 <400> 93
 Tyr Asn Arg Lys Leu Leu Asp Ile Ala
 <210> 94
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<212> PRT
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                         10
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 Thr Leu Leu Glu Leu Ala Arg Thr Gln Ser Gln Arg Glu Arg Ala Glu
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                                25
                                                  30
 Gln Asn Arg Ile Ile Phe Asp Ser Val
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 <211> 41
 <212> PRT
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             5 10
   1
                                       1.5
 Thr Leu Leu Glu Leu Ala Arg Thr Gln Ser Gln Arg Glu Arg Ala Glu
             20
                                25
 Gln Asn Arg Ile Ile Phe Asp Ser Val
          35
                            40
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<210> 96
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<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Asp Asn Pro Pro Leu Ser Ile Asp Leu Thr Phe His Leu Leu Arg Thr
                                   10
Leu Leu Glu Leu Ala Arg Thr Gln Ser Gln Arg Glu Arg Ala Glu Gln
                    25
Asn Arg Ile Ile Phe Asp Ser Val
        35
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<211> 40
<212> PRT
<213> Artificial Sequence
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Asp Asp Pro Pro Leu Ser Ile Asp Leu Thr Phe His Leu Leu Arg Thr
                        10
Leu Leu Glu Leu Ala Arg Thr Gln Ser Gln Arg Glu Arg Ala Glu Gln
            20
                               25
Asn Arg Ile Ile Phe Asp Ser Val
        35
                            40
<210> 98
<211> 41
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
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<400> 98
Asn Asp Asp Pro Pro Ile Ser Ile Asp Leu Thr Phe His Leu Leu Arg
          5
                                    10
Asn Met Ile Glu Met Ala Arg Ile Glu Asn Glu Arg Glu Gln Ala Gly
                                 25
Leu Asn Arg Lys Tyr Leu Asp Glu Val
   35
<210> 99
<211> 12
<212> PRT
<213> Artificial Sequence
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<400> 99
Ala Gly Thr Ala Asp Cys Phe Trp Lys Tyr Cys Val
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<210> 100
<211> 13
<212> PRT
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Ala Gly Asn Leu Ser Glu Cys Phe Trp Lys Tyr Cys Val
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 <210> 101
 <211> 46
 <212> PRT
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Peptide

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                             10
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Gly Ala Leu Glu Ser Leu Ile Glu Glu Glu Thr Gly Gln
<210> 105
<211> 13
<212> PRT
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<400> 105
Asp Ala Glu Asn Leu Ile Asp Ser Phe Gln Glu Ile Val
 1 5 10
<210> 106
<211> 13
<212> PRT
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Asn Thr Glu His Leu Val Asp Ser Phe Gln Glu Met Gly
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<210> 107
<211> 13
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Asp Thr Ser His His Asp Gln Asp His Pro Thr Phe Asp
<210> 108
<212> PRT
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<212> PRT
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Trp Cys Leu Glu Ser Ser Gln Cys Gln Asp Leu Ser Thr Glu Ser Asn
                  5
                                 10
                                                          15
Leu Leu Ala Cys Ile Arg Ala Cys Lys Pro
<210> 110
<211> 10
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Peptide <400> 110 Gln His Trp Ser Tyr Gly Leu Ser Pro Gly 1 5 <210> 111 <211> 44 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 111 Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln Leu Ser Ala Arg Lys Leu Gln Asp Ile Met Ser Arg Gln Gln Gly 25 Glu Ser Asn Gln Glu Arg Gly Ala Arg Ala Arg Leu 35 40 <210> 112 <211> 44 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 112 Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln 10 Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Asn Arg Gln Gln Gly 20 25

<210> 113

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Glu Arg Asn Gln Glu Gln Gly Ala Lys Val Arg Leu

<211> 44

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<212> PRT
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Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln
                                    10
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Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly
             20
                                 25
Glu Arg Asn Gln Glu Gln Gly Ala Arg Vai Arg Leu
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<210> 114
<211> 29
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His Ala Asp Ala Ile Phe Thr Ser Ser Tyr Arg Arg Ile Leu Gly Gln
                                     10
                                                         15
Leu Tyr Ala Arg Lys Leu Leu His Glu Ile Met Asn Arg
             20
<210> 115
<211> 30
<212> PRT
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Val Asp Ser Met Trp Ala Glu Gln Lys Gln Met Glu Leu Glu Ser Ile
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Leu Val Ala Leu Leu Gln Lys His Ser Arg Asn Ser Gln Gly
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<210> 116
<211> 29
<212> PRT
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Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln
                       10
      5
Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg
                              25
            20
<210> 117
<211> 29
<212> PRT
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Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln
Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg
            20
<210> 118
<211> 29
<212> PRT
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<400> 118 Tyr Arg Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln 10 Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg 25 <210> 119 <211> 31 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide His Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln 5 10 Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Ser Arg Gln Gln Gly 25 20 <210> 120 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 120 Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln 1 5 10 15 Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly 25 20 Glu Ser Asn Gln Glu 35 <210> 121 <211> 40 <212> PRT

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Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly
                               25
Glu Ser Asn Gln Glu Arg Gly Ala
        35
<210> 122
<211> 40
<212> PRT
<213> Artificial Sequence
<220>
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      Peptide
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Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln
Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly
            20
                                25
Glu Ser Asn Gln Glu Arg Gly Ala
       35
                            40
<210> 123
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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<400> 123
Glu Gln Gly Glu Ser Asn Gln Glu Arg Gly Ala Arg Ala Arg Leu
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1 10 15 <210> 124 <211> 42 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 124 His Val Asp Ala Ile Phe Thr Thr Asn Tyr Arg Lys Leu Leu Ser Gln 10 5 Leu Tyr Ala Arg Lys Val Ile Gln Asp Ile Met Asn Lys Gln Gly Glu 20 25 Arg Ile Gln Glu Gln Arg Ala Arg Leu Ser 35 <210> 125 <211> 44 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Ile Leu Gly Gln 10 5 Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Asn Arg Gln Gln Gly 20 . 25 Glu Arg Asn Gln Glu Gln Gly Ala Lys Val Arg Leu 40 <210> 126 <211> 43 <212> PRT <213> Artificial Sequence

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Leu Tyr Ala Arg Lys Leu Leu His Glu Ile Met Asn Arg Gln Gln Gly
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                                25
Glu Arg Asn Gln Glu Gln Arg Ser Arg Phe Asn
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<210> 127
<211> 6
<212> PRT
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<400> 127
His Trp Ala Trp Phe Lys
<210> 128
<211> 6
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<400> 128
His Trp Ala Trp Phe Lys
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  1
<210> 129
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<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic
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<400> 129
Ala Ala Trp Phe Lys
<210> 130
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<400> 130
His Trp Lys Trp Phe Lys
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<210> 131
<211> 6
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 131
Ala Ala Trp Phe Leu
 1
<210> 132
<211> 45
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 132
His Ala Asp Gly Met Phe Asn Lys Ala Tyr Arg Lys Ala Leu Gly Gln
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10 15 Leu Ser Ala Arg Lys Tyr Leu His Thr Leu Met Ala Lys Arg Val Gly 20 25 Gly Gly Ser Met Ile Glu Asp Asp Asn Glu Pro Leu Ser 40 <210> 133 <211> 44 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 133 Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln 10 Leu Ser Ala Arg Lys Leu Gln Asp Ile Asn Ser Arg Gln Gln Gly 20 25 Glu Ser Asn Gln Glu Arg Gly Ala Arg Ala Arg Leu <210> 134 <211> 28 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 134 Tyr Ala Asp Ala Ile Phe Thr Asn Cys Tyr Arg Lys Val Leu Cys Gln 10 Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Ser Arg 20 25

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<210> 135 <211> 8

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<212> PRT
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Phe Ser Lys Lys Leu Lys Pro Ala
       5
<210> 136
<211> 10
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     Peptide
<400> 136
Glu His Trp Ser His Gly Trp Tyr Pro Gly
       5
<210> 137
<211> 10
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<400> 137
Glu His Trp Ser Tyr Gly Leu Arg Pro Gly
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     Peptide
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<400> 138
Phe Ser Tyr Leu Arg Pro Ala
            5
<210> 139
<211> 10
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     Peptide
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Glu His Trp Ser Tyr Ala Leu Arg Pro Gly
                5
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<400> 140
Glu His Trp Ser Tyr Gly Leu Gln Pro Gly
1 5
<210> 141
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His Trp Ser Tyr Val Arg Pro
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<210> 142
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 142
Glu His Trp Ser Tyr Lys Leu Arg Pro Gly
      5
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<211> 10
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<400> 143
Glu Phe Pro Ser Tyr Phe Leu Arg Pro Gly
                5
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<400> 144
Phe Trp Ser Tyr Ala Leu Arg Pro Gly
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<400> 145
Glu Phe Trp Ser Tyr Trp Leu Arg Pro Gly
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<400> 146
Glu His Trp Ser Tyr Gly Leu Arg Pro
    5
<210> 147
<211> 9
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     Peptide
<400> 147
Glu His Trp Ser Tyr Ala Leu Arg Pro
                5
<210> 148
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<212> PRT
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     Peptide
<400> 148
His Trp Ser Tyr Trp Leu Arg Pro Gly
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      Peptide
<400> 149
Glu His Trp Ser Tyr Trp Leu Arg Pro
                5
<210> 150
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      Peptide
<400> 150
His Trp Ser Tyr Ser Leu Arg Pro
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<210> 151
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      Peptide
<400> 151
Glu His Trp Ser Tyr Arg Trp Leu Pro
<210> 152
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<212> PRT
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     Peptide
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Leu Arg Pro Gly
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<400> 153
His Trp Ser Tyr Gly Leu Arg Pro Gly
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<400> 154
Glu His Tyr Ser Leu Glu Trp Lys Pro Gly
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<211> 10
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1 5 10
<210> 156
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Ser Tyr Gly Leu Arg Pro Gly
      5
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     Peptide
<400> 157
His Trp Ser Tyr Gly Leu Lys Pro Gly
1 5
<210> 158
<211> 7
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    Peptide
<400> 158
Phe Phe Ser Tyr Leu Arg Pro
<210> 159
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<212> PRT
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      Peptide
<400> 159
His Trp Ser Tyr Gly Trp Leu Pro Gly
                5
<210> 160
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      Peptide
<400> 160
His Trp Ser Tyr Ser Leu Arg Pro Gly
<210> 161
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<212> PRT
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<400> 161
Ala Phe Trp Ser Tyr Ser Leu Arg Pro
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          5
<210> 162
<211> 31
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 162
Glu Pro Asp Cys Cys Arg Gln Lys Thr Cys Ser Cys Arg Leu Tyr Glu
                                    10
Leu Leu His Gly Ala Gly Asn His Ala Ala Gly Ile Leu Thr Leu
             20
                                 25
<210> 163
<211> 28
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      Peptide
<400> 163
Arg Ser Gly Pro Pro Gly Leu Gln Gly Arg Leu Gln Arg Leu Leu Gln
                                    10
Ala Ser Gly Asn His Ala Ala Gly Ile Leu Thr Met
             20
                                 25
<210> 164
<211> 28
<212> PRT
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      Peptide
<400> 164
Arg Pro Gly Pro Pro Gly Leu Gln Gly Arg Leu Gln Arg Leu Leu Gln
                  5
Ala Asn Gly Asn His Ala Ala Gly Ile Leu Thr Met
             20
<210> 165
<211> 31
<212> PRT
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<220>
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<400> 165
Ser Arg Thr His Arg His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
                                     10
Pro Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro Val Gly Arg Phe
                                 25
             20
<210> 166
<211> 20
<212> PRT
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<400> 166
Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro
                                     10
Val Gly Arg Phe
             20
<210> 167
<211> 31
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     Peptide
<400> 167
Ser Arg Ala His Gln His Ser Met Glu Thr Arg Thr Pro Asp Ile Asn
 1
                  5
                                     10
Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro Val Gly Arg Phe
                                 25
             20
<210> 168
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56

<211> 20

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<212> PRT
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 <223> Description of Artificial Sequence: Synthetic
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 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro
                                      10
 Val Gly Arg Phe
              20
 <210> 169
 <211> 31
 <212> PRT
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       Peptide
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 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
                                      10
 Pro Ala Trp Tyr Ala Ser Arg Gly Ile Arg Pro Val Gly Arg Phe
              20
                                  25
 <210> 170
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 <212> PRT
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       Peptide
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 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro
                                      10
 Val Gly Arg Phe
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<210> 171
<211> 5
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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<400> 171
Ala Tyr Trp Lys Phe
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<210> 172
<211> 14
<212> PRT
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<400> 172
Pro Cys Lys Asn Phe Phe Trp Lys Thr Phe Ser Ser Cys Lys
                                     10
<210> 173
<211> 17
<212> PRT
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Asp Arg Met Pro Cys Arg Asn Phe Phe Trp Lys Thr Phe Ser Ser Cys
                  5
                                     10
Lys
<210> 174
<211> 17
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<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
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Asp Arg Met Pro Cys Arg Asn Phe Phe Trp Lys Thr Phe Ser Ser Cys
                       10
Lys
<210> 175
<211> 14
<212> PRT
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     Peptide
<400> 175
Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
<210> 176
<211> 13
<212> PRT
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<400> 176
Ala Gly Cys Lys Asn Phe Phe Lys Thr Phe Thr Ser Cys
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<210> 177
<211> 13
<212> PRT
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<400> 177
Ala Gly Cys Lys Asn Phe Phe Lys Thr Tyr Thr Ser Cys
<210> 178
<211> 11
<212> PRT
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<400> 178
Cys His His Phe Phe Lys Thr Phe Thr Ser Cys
         5
<210> 179
<211> 14
<212> PRT
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<400> 179
Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Trp Thr Ser Cys
 1 5
<210> 180
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<400> 180
Phe Cys Tyr Thr Thr
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1
<210> 181
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<400> 181
Ser Ala Asn Ser Asn Pro Ala Ala Pro Arg Glu Arg Lys Ala Gly Cys
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Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
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<210> 182
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<400> 182
Phe Cys Phe Lys Thr Cys Thr
1 , 5
<210> 183
<211> 32
<212> PRT
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      Peptide
<400> 183
Ala Pro Ser Asp Pro Arg Leu Arg Gln Phe Leu Gln Lys Ser Leu Ala
Ala Ala Ala Gly Lys Gln Glu Leu Ala Lys Tyr Phe Leu Ala Glu Leu
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20 25 30

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<210> 184
<211> 15
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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<400> 184
Tyr Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
               5
                              10
<210> 185
<211> 14
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Tyr Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
                                   10
<210> 186
<211> 14
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Ser Ala Asn Ser Asn Pro Ala Met Ala Pro Arg Tyr Arg Lys
      5
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<210> 187
<211> 14
<212> PRT
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     Peptide
<400> 187
Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Tyr Thr Ser Cys
1 5
<210> 188
<211> 15
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<400> 188
Tyr Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
                                10
            5
<210> 189
<211> 14
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     Peptide
<400> 189
Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
1 5
<210> 190
<211> 4
<212> PRT
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<220>
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<400> 190
Phe Trp Lys Thr
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<210> 191
<211> 25
<212> PRT
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     Peptide
<400> 191
Ser Asn Pro Ala Met Ala Pro Arg Glu Arg Lys Ala Gly Cys Lys Asn
1 5 10
Phe Phe Trp Lys Thr Phe Thr Ser Cys
          20
<210> 192
<211> 28
<212> PRT
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     Peptide
Ser Ala Asn Ser Asn Pro Ala Met Ala Pro Arg Glu Arg Lys Ala Gly
     5 10
                                        15
Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
           20
<210> 193
<211> 12
<212> PRT
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     Peptide
<400> 193
Ser Ala Asn Ser Asn Pro Ala Met Ala Pro Arg Glu
       5
<210> 194
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<212> PRT
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     Peptide
<400> 194
Tyr Ser Ala Asn Ser Asn Pro Ala Met Ala Pro Arg Glu Arg Lys Ala
1
      5
                       10
                                               15
Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
          20
                            25
<210> 195
<211> 28
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 195
Ser Ala Asn Ser Asn Pro Ala Leu Ala Pro Arg Glu Arg Lys Ala Gly
     5
                     10
Cys Lys Asn Phe Phe Trp Lys Thr Tyr Thr Ser Cys
          20 25
<210> 196
<211> 14
<212> PRT
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<220>
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 196
Ser Ala Asn Ser Asn Pro Ala Met Ala Pro Arg Glu Arg Lys
                5
                                   10
<210> 197
<211> 8
<212> PRT
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Phe Cys Tyr Trp Lys Val Cys Trp
    5
<210> 198
<211> 8
<212> PRT
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Phe Cys Tyr Trp Lys Val Cys Trp
<210> 199
<211> 6
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Phe Cys Tyr Trp Thr Thr
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<210> 200
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Ala Cys Tyr Trp Leu Val Cys Thr
<210> 201
<211> 14
<212> PRT
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      Peptide
<400> 201
Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
                 5
                                    10
<210> 202
<211> 3
<212> PRT
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<400> 202
His Pro Gly
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<210> 203
<211> 1
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 203
Thr
 1
<210> 204
<211> 3
<212> PRT
<213> Artificial Sequence
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      Peptide
<400> 204
Thr His Pro
 1
<210> 205
<211> 3
<212> PRT
<213> Artificial Sequence
<220>
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      Peptide
<400> 205
Glu His Pro
 1
<210> 206
<211> 3
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 206
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Glu Gln Pro
 1
<210> 207
<211> 2
<212> PRT
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<220>
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<400> 207
Glu His
1
<210> 208
<211> 4
<212> PRT
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      Peptide
<400> 208
His Pro Gly Lys
 1
<210> 209
<211> 3
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<213> Artificial Sequence
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      Peptide
<400> 209
His Pro Gly
 1
<210> 210
<211> 2
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<212> PRT
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<220>
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      Peptide
<400> 210
Glu Pro
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<210> 211
<211> 2
<212> PRT
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<400> 211
Phe Pro
 1
<210> 212
<211> 22
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 212
Phe Leu Trp Lys Asp Leu Gln Arg Val Arg Gly Asp Leu Gly Ala Ala
                                    10
Leu Asp Ser Trp Ile Thr
             20
<210> 213
<211> 24
<212> PRT
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<220>
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     Peptide
<400> 213
Glu Glu Glu Glu Lys Asp Ile Glu Ala Glu Glu Arg Gly Asp Leu Gly
                                  10
Glu Gly Gly Ala Trp Arg Leu His
           20
<210> 214
<211> 10
<212> PRT
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     Peptide
<400> 214
Ser Phe Pro Trp Met Glu Ser Asp Val Thr
1 5
<210> 215
<211> 32
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 215
Cys Ala Ser Leu Ser Thr Cys Val Leu Gly Lys Leu Ser Gln Glu Leu
                5
                                   10
His Lys Leu Gln Thr Tyr Pro Arg Thr Asp Val Gly Ala Gly Thr Pro
                                25
            20
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<210> 216 <211> 32

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<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
Cys Ser Asn Leu Ser Thr Cys Val Leu Gly Lys Leu Ser Gln Glu Leu
                      10
His Lys Leu Gln Thr Tyr Pro Arg Thr Asp Val Gly Ala Gly Thr Pro
            20
                              25
                                                30
<210> 217
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
Cys Gly Asn Leu Ser Thr Cys Met Leu Gly Thr Tyr Thr Gln Asp Phe
 1
     5
                       10
Asn Lys Phe His Thr Phe Pro Gln Thr Ala Ile Gly Val Gly Ala Pro
            20
                              25
<210> 218
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 218
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Cys Ser Asn Leu Ser Thr Cys Val Leu Ser Ala Tyr Trp Arg Asn Leu 1 5 10 15

Asn Asn Phe His Arg Phe Ser Gly Met Gly Phe Gly Pro Glu Thr Pro $20 \\ 25 \\ 30$

<210> 219

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 219

Cys Gly Asn Leu Ser Thr Cys Met Leu Gly Thr Tyr Thr Gln Asp Leu

1 5 10 15

Asn Lys Phe His Thr Phe Pro Gln Thr Ser Ile Gly Val Gly Ala Pro 20 25 30

<210> 220

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 220

Cys Ser Asn Leu Ser Thr Cys Val Leu Gly Lys Leu Ser Gln Glu Leu 1 5 10 15

His Lys Leu Gln Thr Tyr Pro Arg Thr Asn Thr Gly Ser Gly Thr Pro
20 25 30

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<210> 221
<211> 25
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 221
Val Leu Gly Lys Leu Ser Gln Glu Leu His Lys Leu Gln Thr Tyr Pro
                                     10
Arg Thr Asn Thr Gly Ser Gly Thr Pro
             20
<210> 222
<211> 21
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 222
Asp Met Ser Ser Asp Leu Glu Arg Asp His Arg Pro His Val Ser Met
                  5
                                    10
Pro Gln Asn Ala Asn
             20
<210> 223
<211> 57
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 223
Ala Pro Phe Arg Ser Ala Leu Glu Ser Ser Pro Ala Asp Pro Ala Thr
                  5
                                    10
                                                         15
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Leu Ser Glu Asp Glu Ala Arg Leu Leu Leu Ala Ala Leu Val Gln Asp
             20
                                25
Tyr Val Gln Met Lys Ala Ser Glu Leu Glu Gln Glu Gln Glu Arg Glu
                             40
Gly Ser Ser Leu Asp Ser Pro Arg Ser
     50
                        55
<210> 224
<211> 37
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 224
Ser Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
                5
Ser Arg Ser Gly Gly Val Val Lys Ser Asn Phe Val Pro Thr Asn Val
             20
                                25
Gly Ser Gln Ala Phe
         35
<210> 225
<211> 19
<212> PRT
<213> Artific al Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 225
Ser Gly Gly Val Val Lys Asn Asn Phe Val Pro Thr Asn Val Gly Ser
                  5
                                    10
                                                        15
Lys Ala Phe
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<210> 226
<211> 19
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 226
Ser Gly Gly Val Val Lys Asn Asn Phe Val Pro Thr Asn Val Gly Ser
                          10
Lys Ala P':
<210> 227
<211> 15
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 227
Val Lys Asn Asn Phe Val Pro Thr Asn Val Gly Ser Lys Ala Phe
                                    10
<210> 228
<211> 37
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 228
Ser Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
Ser Arg Ser Gly Gly Val Val Lys Asp Asn Phe Val Pro Thr Asn Val
             20
                                 25
Gly Ser Lys Ala Phe
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<210> 229 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 229 Ala Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Asp Phe Leu 5 Ser Arg Ser Gly Gly Val Gly Lys Asn Asn Phe Val Pro Thr Asn Val Gly Ser Lys Ala Phe 35 <210> 230 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 230 Ala Cys Asp Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu Ser Arg Ser Gly Gly Val Val Lys Asn Asn Phe Val Pro Thr Asn Val 20 . 25 Gly Ser Lys Ala Phe 35 <210> 231 <211> 7 <212> PRT

<213> Artificial Sequence

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<220>
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  Peptide
<400> 231
Cys Gly Asn Leu Ser Thr Cys
 1 5
<210> 232
<211> 16
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 232
Asp Met Ala Lys Asp Leu Glu Thr Asn His Pro Tyr Phe Gly Asn
                 5
                            10
<210> 233
<211> 19
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 233
Ala Cys Asp Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
                5 10
Ser Arg Ser
 <210> 234
 <211> 18
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 234
Gly Gly Val Val Lys Asn Asn Phe Val Pro Thr Asn Val Gly Ser Lys
                                    10
Ala Phe
<210> 235
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 235
Val Thr His Arg Leu Ala Gly Leu Leu Ser Arg Ser Gly Gly Val Val
                  5
                                    10
Lys Asn Asn Phe Val Pro Thr Asn Val Gly Ser Lys Ala Phe
             20
                                 25
<210> 236
<211> 37
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 236
Ala Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
                 5
                                    10
Ser Arg Ser Gly Gly Met Val Lys Ser Asn Phe Val Pro Thr Asn Val
             20
                                 25
                                                      30
Gly Ser Lys Ala Phe
         35
<210> 237
<211> 37
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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 237
Ser Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
                     10
Ser Arg Ser Gly Gly Val Val Lys Asp Asn Phe Val Pro Thr Asn Val
                              25
Gly Ser Glu Ala Phe
        35
<210> 238
<211> 37
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
Ser Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
    5
                                   10
                                                      15
Ser Arg Ser Gly Gly Val Val Lys Asp Asn Phe Val Pro Thr Asn Val
            20
                               25
Gly Ser Glu Ala Phe
         35
<210> 239
<211> 30
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 239
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Val Thr His Arg Leu Ala Gly Leu Leu Ser Arg Ser Gly Gly Val Val
                                  10
Lys Asp Asn Phe Val Pro Thr Asn Val Gly Ser Glu Ala Phe
                  25
<210> 240
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 240
Pro Thr Asn Val Gly Ser Glu Ala Phe
      5
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<210> 241
<211> 8
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 241
Thr Asn Val Gly Ser Glu Ala Phe
<210> 242
<211> 7
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 242
Asn Val Gly Ser Glu Ala Phe
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<210> 243
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 243
Val Gly Ser Glu Ala Phe
                5
<210> 244
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 244
Val Gly Ser Glu Ala Phe
<210> 245
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 245
Ser Glu Ala Phe
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<210> 246
<211> 37
<212> PRT
<213> Artificial Sequence
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<220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 246 Ala Cys Asp Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu 10 Ser Arg Ser Gly Gly Val Val Lys Asn Asn Phe Val Pro Thr Asn Val 25 Gly Ser Lys Ala Phe 35 <210> 247 <211> 30 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 247 Ser Asn Leu Ser Thr Val Leu Gly Lys Leu Ser Gln Glu Leu His Lys 10 15 1 5 Leu Gln Thr Tyr Pro Arg Thr Asp Val Gly Ala Gly Thr Pro 20 25 <210> 248 <211> 38 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 248 Tyr Ala Cys Asp Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu Ser Arg Ser Gly Gly Val Val Lys Asn Asn Phe Val Pro Thr Asn 25

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Val Gly Ser Lys Ala Phe
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<210> 249
<211> 38
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 249
Tyr Ala Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu
                5
                                    10
Leu Ser Arg Ser Gly Gly Met Val Lys Ser Asn Phe Val Pro Thr Asn
                                25
Val Gly Ser Lys Ala Phe
        35
<210> 250
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 250
Tyr Val Pro Thr Asn Val Gly Ser Glu Ala Phe
 1
                 5
                                     10
<210> 251
<211> 38
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 251
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Tyr Ser Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu
                                   10
Leu Ser Arg Ser Gly Gly Val Val Lys Asp Asn Phe Val Pro Thr Asn
                               25
Val Gly Ser Glu Ala Phe
        35
<210> 252
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 252
Tyr Val Lys Asp Asn Phe Val Pro Thr Asn Val Gly Ser Glu Ala Phe
                                   10
<210> 253
<211> 37
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 253
Ser Cys Asn Thr Ala Thr Cys Val Thr His Arg Leu Ala Gly Leu Leu
                                     10
Ser Arg Ser Gly Gly Val Val Lys Asp Asn Phe Val Pro Thr Asn Val
                                 25
Gly Ser Glu Ala Phe
        35
<210> 254
<211> 46
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic

<220>

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Peptide
<400> 254
Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln Arg Pro Arg Lys Lys
               5
Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser Leu Gly Glu Ala
                                25
Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys Ser Gln
        35
                            40
<210> 255
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 255
Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser Leu Gly
                5
Glu Ala Asp Lys Ala Asp Val Asp Val Leu Thr Lys Ala Lys Ser Gln
                                25
            20
<210> 256
<211> 84
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 256
Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
       5
                                   10
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Ser Met Glu Arg Val Glu Trp Leu Arg Lys Leu Gln Asp Val His 25 20 Asn Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser 40 Gln Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu 55 Lys Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys 75 Ala Lys Ser Gln <210> 257 <211> 21 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 257 Glu Lys Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr 10 Lys Ala Lys Ser Gln 20 <210> 258 <211> 34 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 258 Ser Val Ser Glu Ile Gln Leu Asn His Asn Leu Gly Lys His Leu Asn 10

Ser Leu Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His

20 25 30

Asn Phe

<210> 259

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 259

Ser Val Ser Glu Cys Gln Leu Met His Asn Leu Gly Lys His Leu Asn 1 5 10 15

Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Cys Gln Asp Val His
20 25 30

Asn Phe

<210> 260

<211> 40

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 260

Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile Gln

1 5 10 15

Asp Leu Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile His
20 25 30

Thr Ala Glu Ile Arg Ala Thr Ser 35 40

<210> 261

<211> 34

<212> PRT

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<213> Artificial Sequence
    <220>
    <223> Description of Artificial Sequence: Synthetic
          Peptide
    <400> 261
    Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
    Ser Leu Glu Arg Val Glu Trp Leu Arg Lys Leu Gln Asp Val His
                20
                                 25
    Asn Phe
    <210> 262
    <211> 32
    <212> PRT
    <213> Artificial Sequence
    <220>
    <223> Description of Artificial Sequence: Synthetic
         Peptide
    <400> 262
    Ala Val Ser Glu Ile Gln Phe His Asn Leu Gly Lys His Leu Ser Ser
    Glu Arg Val Glu Trp Leu Arg Lys Leu Gln Asp Val His Asn Tyr
                                 25
<210> 263
    <211> 30
    <212> PRT
    <213> Artificial Sequence
    <220>
    <223> Description of Artificial Sequence: Synthetic
         Peptide
    <400> 263
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Ser Glu Ile Gln Phe His Asn Leu Gly Lys His Leu Ser Ser Glu Arg

1 5 10 15

Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn Tyr 20 25 30

<210> 264

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 264

Ser Val Ser Glu Ile Gln Leu His Asn Leu Gly Lys His Leu Asn Ser 1 5 10 15

Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn Tyr
20 25 30

<210> 265

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 265

Ser Val Ser Glu Ile Gln Leu His Asn Leu Gly Lys His Leu Asn Ser 1 5 10 15

Glu Arg Val Glu Trp Leu Arg Lys Leu Gln Asp Val His Asn Tyr
20 25 30

<210> 266

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<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 266
Ser Glu Ile Gln Leu His Asn Leu Gly Lys His Leu Asn Ser Glu Arg
                                   10
Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn Tyr
            20
                               25
<210> 267
<211> 26
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 267
Phe His Asn Leu Gly Lys His Leu Ser Ser Glu Arg Val Glu Trp Leu
                5
Arg Lys Lys Leu Gln Asp Val His Asn Tyr
            20
<210> 268
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
Ala Val Ser Glu Ile Gln Leu His Asn Leu Gly Lys His Leu Ala Ser
                                   10
                5
                                                      15
Val Glu Arg Gln Trp Leu Arg Lys Leu Gln Asp Val His Asn Tyr
            20
                              25
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<211> 25
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 269
Arg Asp Ala Gly Ser Gln Arg Pro Arg Lys Lys Glu Asp Asn Val Leu
                        10
Val Glu Ser His Glu Lys Ser Leu Gly
            20
<210> 270
<211> 32
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 270
Ser Glu Ile Gln Phe Met His Asn Leu Gly Lys His Leu Ser Ser Met
                                  10
Glu Arg Val Glu Trp Leu Arg Lys Leu Gln Asp Val His Asn Phe
            20
                    25
<210> 271
<211> 31
<212> PRT
<213> Artificial Sequence
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<210> 269

<220>

92

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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 271
Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
                                    10
Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val
                                25
<210> 272
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 272
Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His
                                25
             20
Asn Phe
<210> 273
<211> 22
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 273
Lys His Leu Asn Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu
                                    10
Gln Asp Val His Asn Phe
             20
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<210> 274
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 274
Ala Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Ala
                                  10
Ser Val Glu Arg Met Gln Trp Leu Arg Lys Leu Gln Asp Val His
            20
                  25
Asn Phe
<210> 275
<211> 38
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 275
Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
Ser Met Glu Arg Val Glu Trp Leu Arg Lys Leu Gln Asp Val His
                    25
            20
Asn Phe Val Ala Leu Gly
       35
<210> 276
<211> 44
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
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<400> 276 Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn 5 10 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His 25 Asn Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg 35 40 <210> 277 <211> 21 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 277 Leu Gln Asp Val His Asn Phe Val Ala Leu Gly Ala Pro Leu Ala Pro 10 Arg Asp Ala Gly Ser 20 <210> 278 <211> 30 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 278 Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln Arg Pro Arg Lys Lys 10 Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser Leu Gly 20 25 <210> 279 <211> 46

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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 279
Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln Arg Pro Arg Lys Lys
Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser Leu Gly Glu Ala
Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys Ser Gln
                . 40
<210> 280
<211> 32
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser Leu Gly
 1
                5
Glu Ala Asp Lys Ala Asp Val Asp Val Leu Thr Lys Ala Lys Ser Gln
                               25
            20
<210> 281
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 281

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Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys Ser Gln
 1 5
                                   10
<210> 282
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 282
Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys Ser Gln
                                  10
<210> 283
<211> 36
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial' Sequence: Synthetic
     Peptide
Tyr Pro Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu
 1
                 5
                                    10
                                                       15
Leu Asn Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr
            20
                                25
Arg Pro Arg Tyr
       35
<210> 284
<211> 41
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 284
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Tyr Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile

1 5 10 15

Gln Asp Leu Arg Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile 20 25 30

His Thr Ala Glu Ile Arg Ala Thr Ser

<210> 285

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 285

Tyr Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu

1 5 10 15

Asn Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val 20 25 30

His Asn Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg
35 40 45

<210> 286

<211> 35

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Peptide

<400> 286

Tyr Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu

1 5 10 15

Asn Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val 20 25 30

His Asn Phe

35

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<211> 34
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 287
Tyr Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn
                                     10
Ser Met Glu Arg Val Glu Trp Leu Arg Lys Leu Gln Asp Val His
                                 25
Asn Phe
<210> 288
<211> 22
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 288
Tyr Leu Gln Asp Val His Asn Phe Val Ala Leu Gly Ala Pro Leu Ala
                                                         15
Pro Arg Asp Ala Gly Ser
             20
<210> 289
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<210> 287

<400> 289 Phe Met His Asn Leu Gly Lys His Leu Ser Ser Met Glu Arg Val Glu 5 10 15 Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn Tyr 20 <210> 290 <211> 26 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 290 Tyr Arg Asp Ala Gly Ser Gln Arg Pro Arg Lys Lys Glu Asp Asn Val 1 5 Leu Val Glu Ser His Glu Lys Ser Leu Gly 20 <210> 291 <211> 33 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 291 Tyr Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser Leu 5 10 . 15 Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys Ser 25 20 Gln <210> 292 <211> 22

100

<212> PRT

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 292
Tyr Glu Lys Ser Leu Gly Glu Ala Asp Lys Ala Asp Val Asn Val Leu
Thr Lys Ala Lys Ser Gln
            20
<210> 293
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 293
Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile Gln
                                  10
                 5
Asp Leu Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile His
            20
                    25
Thr Ala
<210> 294
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 294
Ala Arg Ser Ala Trp
  1
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<210> 295
<211> 34
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 295
Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile Gln
                                   10
Asp Leu rg Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile His
                               25
            20
Thr Ala
<210> 296
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 296
Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile Gln Asp
                                    10
Leu Arg Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile His Thr
            20
                                25
Ala
<210> 297
<211> 35
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 297

Tyr Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile 1 . 5 10 Gln Asp Leu Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile 25 His Thr Ala 35 <210> 298 <211> 34 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 298 Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile Gln 10 Asp Leu Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile His 25 Thr Tyr <210> 299 <211> 37 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 299 Ala Val Ser Glu His Gln Leu Leu His Asp Lys Gly Lys Ser Ile Gln 10 Asp Leu Arg Arg Phe Phe Leu His His Leu Ile Ala Glu Ile His 20 25

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Thr Ala Glu Ile Arg
        35
<210> 300
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 300
Leu Leu His Asp Lys Gly Lys Ser Ile Gln Asp Leu Arg Arg Phe
Phe Leu His His Leu Ile Ala Glu Ile His Thr Ala
             20
                                25
<210> 301
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 301
Ala Thr Ser Glu Val Ser Pro Asn Ser Lys Pro Ser Pro Asn Thr Lys
                 5
                                    10
                                                         15
Asn His Pro Val Arg Phe Gly Ser Asp Asp Glu
             20
<210> 302
<211> 20
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 302
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Tyr Leu Thr Gln Glu Thr Asn Lys Val Glu Thr Tyr Lys Glu Gln Pro
                                     10
 Leu Lys Thr Pro
             20
 <210> 303
 <211> 5
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 303
 Thr Arg Ser Ala Trp
  1
 <210> 304
 <211> 5
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 304
 Thr Arg Ser Ala Trp
  1
 <210> 305
<211> 32
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 305
 Thr Arg Ser Ala Trp Leu Asp Ser Gly Val Thr Gly Ser Gly Leu Glu
                   5
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Gly Asp His Leu Ser Asp Thr Ser Thr Thr Ser Leu Glu Leu Asp Ser
20 25 30

<210> 306

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 306

Ser Ala Trp Leu Asp Ser Gly Val Thr Gly Ser Gly Leu Glu Gly Asp
1 5 10 15

His Leu Ser Asp Thr Ser Thr Thr Ser Leu Glu Leu Asp Ser Arg Arg 20 25 30

His

<210> 307

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 307

Thr Ala Leu Leu Trp Gly Leu Lys Lys Lys Lys Glu Asn Asn Arg Arg

1 5 10 15

Thr His His Met Gln Leu Met Ile Ser Leu Phe Lys Ser Pro Leu Leu 20 25 30

Leu Leu

<210> 308

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<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 308
Met Ile Pro Ala Lys Asp Met Ala Lys Val Met Ile Val Met Leu Ala
Ile Arg Phe Leu Thr Lys Ser Asp Gly Lys Ser Val Lys Lys Arg
                               25
<210> 309
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 309
Thr Ala Leu Leu Trp Gly Leu Lys Lys Lys Lys Gly Lys Gln Gln Lys
                5
                                  10
Asn Thr Ser Tyr Ala Thr Asn Asp Leu Ile Ile
            20
                               25
<210> 310
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 310
Ala Thr Gln Arg Leu Ala Asn Phe Leu Val His Ser Ser Asn Asn Phe
                        10
Gly Ala Ile Leu Ser Ser Thr Asn Val Gly Ser Asn Thr Tyr
            20
                              25
                                                  30
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<210> 311
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 311
Ala Thr Gln Arg Leu Ala Asn Phe Leu Val Arg Ser Ser Asn Asn Leu
                                     10
Gly Pro Val Leu Pro Pro Thr Asn Val Gly Ser Asn Thr Tyr
                                 25
<210> 312
<211> 25
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 312
Val Leu Gly Lys Leu Ser Gln Glu Leu His Lys Leu Gln Thr Tyr Pro
                  5
                                                         15
Arg Thr Asn Thr Gly Ser Asn Thr Tyr
             20
                                 25
<210> 313
<211> 24
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 313
Leu Gly Arg Leu Ser Gln Glu Leu His Arg Leu Gln Thr Tyr Pro Arg
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                  5
                                     10
                                                          15
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Thr Asn Thr Gly Ser Asn Thr Tyr
              20
 <210> 314
 <211> 25
 <212> PRT
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 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 314
 Ala Thr Gln Arg Leu Ala Asn Glu Leu Val Arg Leu Gln Thr Tyr Pro
                                     10
                                                          15
 Arg Thr Asn Val Gly Ser Asn Thr Tyr
              20
<210> 315
 <211> 29
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 315
 Thr Gln Arg Leu Ala Asn Phe Leu Val His Ser Ser Asn Asn Phe Gly
 Ala Ile Leu Ser Ser Thr Asn Val Gly Ser Asn Thr Tyr
             20 25
 <210> 316
  <211> 37
  <212> PRT
  <213> Artificial Sequence
 <220>
  <223> Description of Artificial Sequence: Synthetic
       Peptide
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<400> 316 Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu 5 10 Ile Arg Ser Ser Asn Asn Leu Gly Ala Ile Leu Ser Pro Thr Asn Val 25 Gly Ser Asn Thr Tyr 35 <210> 317 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu 10 Val His Ser Ser Asn Asn Phe Gly Ala Ile Leu Ser Ser Thr Asn Val 20 25 Gly Ser Asn Thr Tyr 35 <210> 318 <211> 37 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu 5 10 Val His Ser Ser Asn Asn Phe Gly Ala Ile Leu Ser Ser Thr Asn Val 20 25 30

Gly Ser Asn Thr Tyr

35

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<210> 319
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala
<210> 320
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 320
Ser Asn Asn Phe Gly Ala Ile Leu Ser Ser
       5
<210> 321
<211> 37
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu
 1
           5
                                   10
Val Arg Ser Ser Asn Asn Leu Gly Pro Val Leu Pro Pro Thr Asn Val
                                25
                                                    30
             20
Gly Ser Asn Thr Tyr
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35

<210> 322 <211> 30 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 322 Ala Thr Gln Arg Leu Ala Asn Phe Leu Val Arg Ser Ser Asn Asn Leu 5 10 Gly Pro Val Leu Pro Pro Thr Asn Val Gly Ser Asn Thr Tyr 20 25 <210> 323 <211> 30 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 323 Ala Thr Gln Arg Leu Ala Asn Phe Leu Val His Ser Ser Asn Asn Phe 10 Gly Ala Ile Leu Ser Ser Thr Asn Val Gly Ser Asn Thr Tyr 25 20 <210> 324 <211> 30 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 324

112

Ala Thr Gln Arg Leu Ala Asn Phe Leu Val Arg Ser Ser Asn Asn Leu

10 15 1 5 Gly Pro Val Leu Pro Pro Thr Asn Val Gly Ser Asn Thr Tyr 20 25 <210> 325 <211> 25 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide Val Leu Gly Lys Leu Ser Gln Glu Leu His Lys Leu Gln Thr Tyr Pro 10 Arg Thr Asn Thr Gly Ser Asn Thr Tyr 20 <210> 326 <211> 24 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 326 Leu Gly Arg Leu Ser Gln Glu Leu His Arg Leu Gln Thr Tyr Pro Arg 5 Thr Asn Thr Gly Ser Asn Thr Tyr 20 <210> 327 <211> 25 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide

<400> 327 Ala Thr Gln Arg Leu Ala Asn Glu Leu Val Arg Leu Gln Thr Tyr Pro 10 Arg Thr Asn Val Gly Ser Asn Thr Tyr 20 <210> 328 <211> 29 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 328 Thr Gln Arg Leu Ala Asn Phe Leu Val His Ser Ser Asn Asn Phe Gly 5 10 Ala Ile Leu Ser Ser Thr Asn Val Gly Ser Asn Thr Tyr 20 <210> 329 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 329 Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu 1 5 10 Ile Arg Ser Ser Asn Asn Leu Gly Ala Ile Leu Ser Pro Thr Asn Val 25 30 Gly Ser Asn Thr Tyr 35 <210> 330

<211> 37

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<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu
Val His Ser Ser Asn Asn Phe Gly Ala Ile Leu Ser Ser Thr Asn Val
                               25
            20
Gly Ser Asn Thr Tyr
     . 35
<210> 331
<211> 37
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 331
Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu
1 5
Val His Ser Ser Asn Asn Phe Gly Ala Ile Leu Ser Ser Thr Asn Val
                               25
            20
Gly Ser Asn Thr Tyr
       35
<210> 332
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 332
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Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala
1 5
<210> 333
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 333
Ser Asn Asn Phe Gly Ala Ile Leu Ser Ser
     5
<210> 334
<211> 37
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
Lys Cys Asn Thr Ala Thr Cys Ala Thr Gln Arg Leu Ala Asn Phe Leu
 1
       5
                                  10
                                                    15
Val Arg Ser Ser Asn Asn Leu Gly Pro Val Leu Pro Pro Thr Asn Val
           20
                             25
Gly Ser Asn Thr Tyr
       35
<210> 335
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 335
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Ala Thr Gln Arg Leu Ala Asn Phe Leu Val Arg Ser Ser Asn Asn Leu 10 Gly Pro Val Leu Pro Pro Thr Asn Val Gly Ser Asn Thr Tyr 20 25 30 <210> 336 <211> 28 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 336 Ser Gln Gly Thr Phe Thr Ser Glu Tyr Ser Lys Tyr Leu Asp Ser Arg 10 Arg Ala Gln Asp Phe Val Gln Trp Leu Met Asn Thr <210> 337 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 337 His Gly Glu Gly Thr Ser Asp Leu Ser Lys Gln Met Glu Glu Ala Arg 5 10 Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser Ser Gly Ala Pro 25 Pro Pro Ser 35 <210> 338 <211> 29 <212> PRT <213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 338
His Ser Gln Gly Thr Phe Thr Ser Asp Tyr Ser Lys Tyr Leu Asp Ser
                  5
                                    10
                                                          15
Arg Arg Ala Gln Asp Phe Val Gln Trp Leu Met Asn Thr
             20
                                 25
<210> 339
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 339
Ala Gln Asp Phe Val Gln Trp Leu Met Asn Thr
                                     10
                  5
<210> 340
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 340
Phe Val Gln Trp Leu Met Asn Thr
<210> 341
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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Peptide

<400> 341

Ser Gln Gly Thr Phe Thr Ser Glu Tyr Ser Lys Tyr Leu Asp Ser Arg

1 5 10 15

Arg Ala Gln Asp Phe Val Gln Trp Leu Met Asn Thr 20 25

<210> 342

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 342

His Asp Glu Phe Glu Arg His Ala Glu Gly Thr Phe Thr Ser Asp Val 1 5 10 15

Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu 20 25 30

Val Lys Gly Arg

<210> 343

<211> 37

<212> PRT

<213> Artificial Sequence

<220>

<400> 343

His Asp Glu Phe Glu Arg His Ala Glu Gly Thr Phe Thr Ser Asp Val $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu $20 \hspace{1cm} 25 \hspace{1cm} 30$

Val Lys Gly Arg Gly

35

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<210> 344
<211> 30
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 344
His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
                                     10
Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
                                 25
<210> 345
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 345
His Ala Asp Gly Ser Phe Ser Asp Glu Met Asn Thr Ile Leu Asp Asn
                                     10
Leu Ala Thr Arg Asp Phe Ile Asn Trp Leu Ile Gln Thr Lys Ile Thr
                                 25
Asp
<210> 346
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 346 His Ala Asp Gly Ser Phe Ser Asp Glu Met Asn Thr Ile Leu Asp Asn 10 5 Leu Ala Ala Arg Asp Phe Ile Asn Trp Leu Ile Gln Thr Lys Ile Thr 25 Asp Arg <210> 347 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 347 His Ser Glu Gly Thr Phe Thr Ser Asp Tyr Ser Lys Tyr Leu Asp Ser 10 Arg Arg Ala Glu Asp Phe Val Glu Trp Leu Met Asn Thr Lys Arg Asn 25 Lys Asn Asn Ile Ala 35 <210> 348 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 348 His Ser Glu Gly Thr Phe Thr Ser Asp Tyr Ser Lys Tyr Leu Asp Ser 5 10 Arg Arg Ala Glu Asp Phe Val Glu Trp Leu Met Asn Thr Lys Arg Asn 25

Lys Asn Asn Ile Ala

35

<210> 349 <211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 349

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser 20 25 30

Ser Gly Ala Pro Pro Pro Ser 35

<210> 350

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 350

His Ala Asp Gly Thr Leu Thr Ser Asp Ile Ser Ser Phe Leu Glu Lys
1 5 10 15

Gln Ala Thr Lys Glu Phe Ile Ala Trp Leu Val Ser Gly Arg Gly Arg 20 25 30

Arg Gln

<210> 351

<211> 29

<212> PRT

<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 351
His Ser Gln Gly Thr Phe Thr Ser Asp Tyr Ser Lys Tyr Leu Asp Ser
 1 5 10
Lys Lys Ala Gln Glu Phe Val Gln Trp Leu Met Asn Thr
           20
                              25
<210> 352
<211> 39
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 352
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
                                  10
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
                              25
Ser Gly Ala Pro Pro Pro Ser
        35
<210> 353
<211> 33
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 353
His Ala Asp Gly Ser Phe Thr Ser Asp Ile Asn Lys Val Leu Asp Thr
                                   10
Ile Ala Ala Lys Glu Phe Leu Asn Trp Leu Ile Ser Thr Lys Val Thr
                               25
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Glu

<220>

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<210> 354
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 354
His Asp Glu Phe Glu Arg His Ala Glu Gly Thr Phe Thr Ser Asp Val
                                     10
Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu
             20
                                 25
Val Lys Gly Arg
         35
<210> 355
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 355
His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
                                     10
Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
             20
                                 25
<210> 356
<211> 19
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic

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Peptide
<400> 356
Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val
                                  10
Xaa Gly Arg
<210> 357
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 357
Ser Asp Val Ser
<210> 358
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 358
Thr Ser Asp Val Ser
           <210> 359
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
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<400> 359

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Phe Thr Ser Asp Val Ser
. 1 5
 <210> 360
 <211> 7
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
      Peptide
 <400> 360
 Thr Phe Thr Ser Asp Val Ser
      5
 <210> 361
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Synthetic
      Peptide
 <400> 361
 Gly Thr Phe Thr Ser Asp Val Ser
 1
      5
 <210> 362
 <211> 9
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
      Peptide
 <400> 362
 Glu Gly Thr Phe Thr Ser Asp Val Ser
 <210> 363
 <211> 10
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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 363
Ala Glu Gly Thr Phe Thr Ser Asp Val Ser
      5
<210> 364
<211> 39
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 364
His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
                                  10
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
                              25
Ser Gly Ala Pro Pro Pro Ser
       35
<210> 365
<211> 39
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
   Peptide
<400> 365
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
                                   10
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
                                25
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Ser Gly Ala Pro Pro Pro Ser
  35
<210> 366
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 366
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
                                   10
Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Tyr
            20
                               25
<210> 367 .
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 367
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
                5
                                    10
Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Tyr
<210> 368
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 368
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Asp Leu Ser Lys Gln Met Glu Glu Glu Ala Val Arg Leu Met Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser Ser Gly Ala Pro Pro Pro Ser 25 <210> 369 <211> 37 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 369 His Asp Glu Phe Glu Arg His Ala Glu Gly Thr Phe Thr Ser Asp Val 5 10 Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu 25 Val Lys Gly Arg Lys 35 <210> 370 <211> 31 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 370 His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly 10 Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg Lys 25 <210> 371 <211> 40 <212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide <400> 371 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu 5 10 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser 20 25 Ser Gly Ala Pro Pro Pro Ser Lys 35 40 <210> 372 <211> 40 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 372 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu 10 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser 20 25 Ser Gly Ala Pro Pro Pro Ser Lys 35 <210> 373 <211> 40 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 373 His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu 5 10 15

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Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
            20
                               25
                                                    30
Ser Gly Ala Pro Pro Pro Ser Lys
        35
<210> 374
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 374
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Glu Met Glu Glu
       5
Glu Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Tyr
            20
                                25
                                                   30
<210> 375
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 375
His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Glu Met Glu Glu
       . 5
Glu Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Tyr
<210> 376
<211> 29
<212> PRT
<213> Artificial Sequence
<220>
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131

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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 376
Asp Leu Ser Lys Gln Met Glu Glu Glu Ala Val Arg Leu Phe Ile Glu
                                   10
Trp Leu Lys Gly Gly Pro Ser Ser Gly Pro Pro Pro Ser
            20
                              25
<210> 377
<211> 31
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 377
Glu Ala Glu Asp Leu Gln Val Gly Gln Val Glu Leu Gly Gly Pro
               5
Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln
                              25
<210> 378
<211> 32
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
Tyr Glu Ala Glu Asp Leu Gln Val Gly Gln Val Glu Leu Gly Gly Gly
                5
                                 10
                                                       15
Pro Gly Ala Gly Ser Leu Gln Pro Leu Ala Leu Glu Gly Ser Leu Gln
                              25
            20
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<210> 379
<211> 51
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 379
Gly Ile Val Glu Gln Cys Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu
                                   10
Glu Asn Tyr Cys Asn Phe Val Asn Gln His Leu Cys Gly Ser His Leu
                       25
           20
Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe Phe Tyr Thr
                            40
Pro Lys Thr
   50
<210> 380
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
Asp Val Ser Thr Pro Pro Thr Val Leu Pro Asp Asn Phe Pro Arg Tyr
                5
                                   10
                                                        15
<210> 381
<211> 35
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 381
Arg Asp Val Ser Thr Pro Pro Thr Val Leu Pro Asp Asn Phe Pro Arg
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1 5 10 15 Tyr Pro Val Gly Lys Phe Phe Gln Tyr Asp Thr Trp Lys Gln Ser Thr 25 Gln Arg Leu 35 <210> 382 <211> 24 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 382 Gly Leu Pro Ala Leu Leu Arg Ala Arg Arg Gly His Val Leu Ala Lys 10 Glu Leu Glu Ala Phe Arg Glu Ala 20 <210> 383 <211> 36 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 383 Tyr Pro Ser Lys Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp 5 10 Met Ala Arg Tyr Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Leu Thr 20 25 Arg Pro Arg Tyr 35 <210> 384 <211> 39

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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 384
Pro Asp Lys Asp Phe Ile Val Asn Pro Ser Asp Leu Val Leu Asp Asn
                         10
Lys Ala Ala Leu Arg Asp Tyr Leu Arg Gln Ile Asn Glu Tyr Phe Ala
                            25
Ile Ile Gly Arg Pro Arg Phe
        35
<210> 385
<211> 3
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 385
Phe Met Arg
 1
<210> 386
<211> 8
<212> PRT.
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
   Peptide
<400> 386
Cys Trp Arg Tyr Cys Trp Arg Tyr
<210> 387
<211> 36
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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 387
Tyr Pro Ser Lys Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp
Met Ala Arg Tyr Tyr Ser Ala Leu Arg His Tyr Ile Asn Leu Ile Thr
               25
           20
Arg Gln Arg Tyr
       35
<210> 388
<211> 24
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 388
Tyr Pro Ser Lys Pro Asp Asn Pro Gly Glu Asp Ala Pro Ala Glu Asp
1 5 10
Met Ala Arg Tyr Tyr Ser Ala Leu
           20
<210> 389
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 389
Ile Gly Pro Tyr Arg Leu Arg Tyr
1 5
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<210> 390
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 390
Gly Pro Ser Gln Pro Thr Tyr Pro Gly Asp Asp Ala Pro Val Glu Asp
Leu Ile Arg Phe Tyr Asp Asn Leu Gln Gln Tyr Leu Asn Val Val Thr
Arg His Arg Tyr
        35
<210> 391
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
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      Peptide
<400> 391
Ala Pro Leu Glu Pro Val Tyr Pro Gly Asp Asn Ala Thr Pro Glu Gln
                 5
                                     10
Met Ala Gln Tyr Ala Ala Asp Leu Arg Arg Tyr Ile Asn Met Leu Thr
                                 25
Arg Pro Arg Tyr
         35
<210> 392
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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Peptide
<400> 392
Leu Thr Arg Pro Arg Tyr
<210> 393
<211> 6
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 393
Leu Thr Arg Pro Arg Tyr
1
         5
<210> 394
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 394
Ala Pro Ser Glu Pro His His Pro Gly Asp Gln Ala Thr Gln Asp Gln
                 5
                                    10
Leu Ala Gln Tyr Tyr Ser Asp Leu Tyr Gln Tyr Ile Thr Phe Val Thr
            20
                               25
Arg Pro Arg Phe
        35
<210> 395
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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Peptide

<400> 395

Ala Pro Leu Glu Pro Met Tyr Pro Gly Asp Tyr Ala Thr His Glu Gln
1 5 10 15

Arg Ala Gln Tyr Glu Thr Gln Leu Arg Arg Tyr Ile Asn Thr Leu Thr 20 25 30

Arg Pro Arg Tyr 35

<210> 396

<211> 36

<212> PRT

<213> Artificial Sequence

<220:

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 396

Tyr Pro Pro Lys Pro Glu Asn Pro Gly Glu Asp Ala Pro Pro Glu Glu

1 5 10 15

Leu Ala Lys Tyr Tyr Thr Ala Leu Arg His Tyr Ile Asn Leu Ile Thr
20 25 30

Arg Gln Arg Tyr 35

<210> 397

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Peptide

<400> 397

Tyr Pro Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu

1 5 10 15

Leu Asn Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr 20 25 30

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Arg Gln Arg Tyr
        35
<210> 398
<211> 34
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 398
Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu Leu Asn
                                   10
Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr Arg Gln
           20
                      25
Arg Tyr
<210> 399
<211> 36
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 399
Tyr Pro Ala Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu
                5
                                   10
Leu Ser Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Val Thr
            20
                                25
Arg Gln Arg Tyr
         35
<210> 400
<211> 36
<212> PRT
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 400
Tyr Pro Ile Lys Pro Glu Ala Pro Gly Glu Asp Ala Ser Pro Glu Glu
                                     10
                  5
Leu Asn Arg Tyr Tyr Ala Ser Leu Arg His Tyr Leu Asn Leu Leu Thr
                                 25
Arg Pro Arg Tyr
        35
<210> 401
<211> 10
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 401
Glu Gln Asp Tyr Thr Gly Trp Met Asp Phe
  1
                  5
<210> 402
<211> 33
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 402
Lys Ala Pro Ser Gly Arg Met Ser Ile Val Lys Asn Leu Gln Asn Leu
                  5
                                      10
Asp Pro Ser His Arg Ile Ser Asp Arg Asp Tyr Met Gly Trp Met Asp
              20
                                  25
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Phe

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<210> 403
   <211> 4
   <212> PRT
   <213> Artificial Sequence
   <220>
   <223> Description of Artificial Sequence: Synthetic
         Peptide
   <400> 403
  Asp Tyr Met Gly
   <210> 404
   <211> 8
   <212> PRT
   <213> Artificial Sequence
   <220>
   <223> Description of Artificial Sequence: Synthetic
         Peptide
   <400> 404
   Asp Tyr Met Gly Trp Met Asp Phe
   1
                   5
   <210> 405
   <211> 8
   <212> PRT
   <213> Artificial Sequence
<220>
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         Peptide
   <400> 405
   Asp Tyr Met Gly Trp Met Asp Phe
    1
                    5
   <210> 406
   <211> 7
   <212> PRT
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     Peptide
<400> 406
Tyr Met Gly Trp Met Asp Phe
1
<210> 407
<211> 4
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     Peptide
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Trp Met Asp Phe
 1
<210> 408
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 408
Lys Ala Pro Ser Gly Arg Val Ser Met Ile Lys Asn Leu Gln Ser Leu
                       10
Asp Pro Ser His Arg Ile Ser Asp Arg Asp Tyr Met Gly Trp Met Asp
           20
                               25
Phe
<210> 409
<212> PRT
<213> Artificial Sequence
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<220>
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<211> 9
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<223> Description of Artificial Sequence: Synthetic
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<400> 410
Ser Ala Glu Glu Tyr Glu Tyr Pro Ser
                5
<210> 411
<211> 5
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      Peptide
<400> 411
Asp Tyr Met Gly Trp
 1
<210> 412
<211> 6
<212> PRT
<213> Artificial Sequence
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      Peptide
<400> 412
Asp Tyr Met Gly Trp Met
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<210> 413
<211> 9
<212> PRT
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<400> 413
Val Pro Val Glu Ala Val Asp Pro Met
<210> 414
<212> PRT
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      Peptide
<400> 414
<210> 415
<211> 8
<212> PRT
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     Peptide
<400> 415
Asp Tyr Met Gly Trp Met Asp Phe
<210> 416
<211> 23
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 416
Met Lys Val Ala Ile Ile Phe Leu Leu Ser Ala Leu Ala Leu Leu Asn
                            10
Leu Ala Gly Asn Thr Thr Ala
           20
<210> 417
<211> 27
<212> PRT
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     Peptide
<400> 417
Val Pro Leu Pro Ala Gly Gly Gly Thr Val Leu Thr Lys Met Tyr Pro
                                    10
                                                       15
Arg Gly Asn His Trp Ala Val Gly His Leu Met
            20
                                25
<210> 418
<211> 16
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 418
Val Pro Leu Pro Ala Gly Gly Gly Thr Val Leu Thr Lys Met Tyr Pro
                                    10
<210> 419
<211> 27
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 419
Ala Pro Val Ser Val Gly Gly Thr Val Leu Ala Lys Met Tyr Pro
                                   10
Arg Gly Asn His Trp Ala Val Gly His Leu Met
            20
<210> 420
<211> 14
<212> PRT
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      Peptide
<400> 420
Met Tyr Pro Arg Gly Asn His Trp Ala Val Gly His Leu Met
                5
<210> 421
<211> 36
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 421
Phe Leu Pro His Val Phe Ala Glu Leu Ser Asp Arg Lys Gly Phe Val
                 5
                                    10
Gln Gly Asn Gly Ala Val Glu Ala Leu His Asp His Phe Tyr Pro Asp
             20
                                 25
                                                     30
Trp Met Asp Phe
        35
<210> 422
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<211> 17

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<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 422
Glu Gly Pro Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met Asp
                                    10
Phe
<210> 423
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 423
Glu Leu Gly Pro Gln Gly Pro Pro His Leu Val Ala Asp Pro Ser Lys
Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met
Asp Phe
<210> 424
<211> 17
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 424
Glu Arg Pro Pro Met Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met Asp
                                    10
                 5
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Phe

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<210> 425
<211> 5
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 425
Ala Trp Met Asp Phe
1
<210> 426
<211> 42
<212> PRT
<213> Artificial Sequence
<220> .
<223> Description of Artificial Sequence: Synthetic
      Peptide
Tyr Ala Glu Gly Thr Phe Ile Ser Asp Tyr Ser Ile Ala Met Asp Lys
 1
                  5
                                     10
                                                          15
Ile His Gln Gln Asp Phe Val Asn Trp Leu Leu Ala Gln Lys Gly Lys
             20
                                 25
                                                      30
Lys Asn Asp Trp Lys His Asn Ile Thr Gln
         35
                             40
<210> 427
<211> 42
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 427
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Tyr Ala Glu Gly Thr Phe Ile Ser Asp Tyr Ser Ile Ala Met Asp Lys 10 Ile Arg Gln Gln Asp Phe Val Asn Trp Leu Leu Ala Gln Lys Gly Lys 25 Lys Ser Asp Trp Lys His Asn Ile Thr Gln 35 <210> 428 <211> 30 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 428 Tyr Ala Glu Gly Thr Phe Ile Ser Asp Tyr Ser Ile Ala Met Asp Lys 10 Ile Arg Gln Gln Asp Phe Val Asn Trp Leu Leu Ala Gln Lys 25 20 <210> 429 <211> 30 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 429 Tyr Ala Glu Gly Thr Phe Ile Ser Asp Tyr Ser Ile Ala Met Asp Lys 10 Ile Arg Gln Gln Asp Phe Val Asn Trp Leu Leu Ala Gln Lys <210> 430 <211> 22 <212> PRT

<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic
    Peptide
<400> 430
Phe Val Pro Ile Phe Thr His Ser Glu Leu Gln Lys Ile Arg Glu Lys
Glu Arg Asn Lys Gly Gln
           20
<210> 431
<211> 22
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
Phe Val Pro Ile Phe Thr Tyr Gly Glu Leu Gln Arg Met Gln Glu Lys
Glu Arg Asn Lys Gly Gln
           20
<210> 432
<211> 22
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
    Peptide
<400> 432
Phe Val Pro Ile Phe Thr Tyr Gly Glu Leu Gln Arg Leu Gln Glu Lys
                         10
Glu Arg Asn Lys Gly Gln
           20
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<210> 433

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<211> 21
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 433
Phe Val Pro Ile Phe Thr Tyr Gly Glu Leu Arg Leu Gln Glu Lys Glu
                                  10
Arg Asn Lys Gly Gln
            20
<210> 434
<211> 9
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 434
Ile Ala Arg Arg His Pro Tyr Phe Leu
<210> 435
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 435
His Ser Asp Gly Thr Phe Thr Ser Glu Leu Ser Arg Leu Arg Glu Ser
        5
                                  10
Ala Arg Leu Gln Arg Leu Leu Gln Gly Leu Val
            20
                   25
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152

<210> 436

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<211> 27
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 436
His Ser Asp Gly Leu Phe Thr Ser Glu Tyr Ser Lys Met Arg Gly Asn
                                     10
Ala Gln Val Gln Lys Phe Ile Gln Asn Leu Met
             20
                                 25
<210> 437
<211> 27
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 437
His Ser Asp Gly Thr Phe Thr Ser Glu Leu Ser Arg Leu Arg Glu Gly
                                     10
                                                         15
Ala Arg Leu Gln Arg Leu Leu Gln Gly Leu Val
             20
                                 25
<210> 438
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
His Ser Asp Gly Thr Phe Thr Ser Glu Leu Ser Arg Leu Arg Asp Ser
                  5
                                                        15
Ala Arg Leu Gln Arg Leu Leu Gln Gly Leu Val
                                25
            20
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<210> 439
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 439
His Ser Asp Gly Thr Phe Thr Ser Glu Leu Ser Arg Leu Gln Asp Ser
                                     10
                  5
Ala Arg Leu Gln Arg Leu Leu Gln Gly Leu Val
            20
<210> 440
<211> 26
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 440
Ser Asp Gly Thr Phe Thr Ser Glu Leu Ser Arg Leu Arg Asp Ser Ala
                  5
Arg Leu Gln Arg Leu Leu Gly Gly Leu Val
           20
<210> 441
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 441
Ala Ala Met Leu Ala Ser Gln Thr Glu Ala Phe Val Pro Ile Phe Thr
                  5
                              10
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Tyr Gly Glu Leu Gln Arg Met Gln Glu Lys Glu Arg Asn Lys Gly Gln
20 25 30

<210> 442

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 442

His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Arg Gln
1 5 10 15

Leu Ala Val Arg Arg Tyr Leu Asn Ser Ile Leu Asn
20 25

<210> 443

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 443

His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Arg Gln
1 5 10 15

Leu Ala Val Arg Arg Tyr Leu Asn Ser Ile Leu Asn Gly Lys Arg $20 \hspace{1.5cm} 25 \hspace{1.5cm} 30$

<210> 444

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

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     Peptide
<400> 444
Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn
               5
<210> 445
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 445
His Ala Asp Gly Val Phe Thr Ser Asp Phe Ser Arg Leu Leu Gly Gln
1 5
                      10
Leu Ser Ala Lys Lys Tyr Leu Glu Ser Leu Ile
            20
<210> 446
<211> 27
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
His Ala Asp Gly Val Phe Thr Ser Asp Tyr Ser Arg Leu Leu Gly Gln
                      10
Ile Ser Ala Lys Lys Tyr Leu Glu Ser Leu Ile
            20
<210> 447
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 447
His Ala Asp Gly Val Phe Thr Ser Asp Phe Ser Lys Leu Leu Gly Gln
                                    10
Leu Ser Ala Lys Lys Tyr Leu Glu Ser Leu Met
             20
                                25
<210> 448
<211> 42
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 448
His Ala Asp Gly Val Phe Thr Ser Asp Phe Ser Lys Leu Leu Gly Gln
                 5
Leu Ser Ala Lys Lys Tyr Leu Glu Ser Leu Met Gly Lys Arg Val Ser
             20
                                25
Ser Asn Ile Ser Glu Asp Pro Val Pro Val
         35
                             40
<210> 449
<211> 12
<212> PRT
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· <220>
 <223> Description of Artificial Sequence: Synthetic
      Peptide
 <400> 449
Val Ser Ser Asn Ile Ser Glu Asp Pro Val Pro Val
 <210> 450
 <211> 15
 <212> PRT
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 450
Ser Ser Glu Gly Glu Ser Pro Asp Phe Pro Glu Glu Leu Glu Lys
                                   10
                5
<210> 451
<211> 21
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 451
Cys Ser Cys Asn Ser Trp Leu Asp Lys Glu Cys Val Tyr Phe Cys His
Leu Asp Ile Ile Trp
            20
<210> 452
<211> 28
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 452
His Ser Asp Ala Val Phe Thr Asp Asn Tyr Ser Arg Phe Arg Lys Gln
                        10
                 5
Met Ala Val Lys Lys Tyr Leu Asn Ser Val Leu Thr
             20
<210> 453
<211> 28
 <212> PRT
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 453
His Ser Asp Ala Leu Phe Thr Asp Thr Tyr Thr Arg Leu Arg Lys Gln
                                    10
Met Ala Met Lys Lys Tyr Leu Asn Ser Val Leu Asn
                                25
             20
<210> 454
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 454
His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
                                    10
Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn
            20
                             25
<210> 455
<211> 28
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 455
His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
                                    10
                                                        15
                  5
Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn
             20
                                 25
```

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<210> 456
<211> 12
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 456
His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg
       5
<210> 457
<211> 19
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 457
Tyr Thr Arg Leu Arg Lys Gln Met Ala Val Lys Lys Tyr Leu Asn Ser
                                   10
Ile Leu Asn
<210> 458
<211> 18
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
Thr Arg Leu Arg Lys Gln Met Ala Val Lys Lys Tyr Leu Asn Ser Ile
                5
                                   10
Leu Asn
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160

<210> 459

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<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 459
His Ser Asp Ala Val Phe Thr Asp Asn Ser Arg Phe Arg Lys Gln Met
Ala Ala Lys Lys Tyr Leu Asn Ser Val Leu Ala
            20
<210> 460
<211> 23
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 460
Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln Met Ala Val Lys Lys
                 5
                                    10
                                                         15
Tyr Leu Asn Ser Ile Leu Asn
             20
<210> 461
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 461
Lys Pro Arg Arg Pro Tyr Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
                 5
                                     10
Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn
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20 25

<210> 462

<211> 29

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 462

Tyr Phe Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln
1 5 10 15

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg 20 25

<210> 463

<211> 28

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 463

His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
1 5 10 15

Leu Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn 20 25

<210> 464

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 464

Leu Met Tyr Pro Thr Tyr Leu Lys

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1
<210> 465
<211> 29
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 465
Met Met Arg Asp Ser Gly Cys Phe Gly Arg Arg Ile Asp Arg Ile Gly
               5
Ser Leu Ser Gly Met Gly Cys Asn Gly Ser Arg Lys Asn
           20
<210> 466
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
Gly Phe Ile Trp Gly Asn Ile Phe Gly His Tyr Ser Gly Asp Phe
                      10
1 5
<210> 467
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 467
Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg
               5 10
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<210> 468
<211> 24
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 468
Ser Ser Asp Cys Phe Gly Ser Arg Ile Asp Arg Ile Gly Ala Gln Ser
                                     10
Gly Met Gly Cys Gly Arg Arg Phe
            20
<210> 469
<211> 20
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 469
Cys Phe Gly Ser Arg Ile Asp Arg Ile Gly Ala Gln Ser Gly Met Gly
                                     10
                                                          15
Cys Gly Arg Phe
             20
<210> 470
<211> 21
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 470
Cys Phe Gly Ser Arg Ile Asp Arg Ile Gly Ala Gln Ser Gly Met Gly
                                      10
                                                          15
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Cys Gly Arg Arg Phe
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20

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<210> 471
<211> 24
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 471
Ser Ser Asp Cys Phe Gly Ser Arg Ile Asp Arg Ile Gly Ala Gln Ser
                                   10
       5
Gly Met Gly Cys Gly Arg Arg Phe
            20
<210> 472
<211> 30
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 472
Ala Pro Arg Ser Met Arg Arg Ser Ser Asp Cys Phe Gly Ser Arg Ile
                                   10
Asp Arg Ile Gly Ala Gln Ser Gly Met Gly Cys Gly Arg Phe
             20
<210> 473
<211> 24
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 473
Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly Ala Gln Ser Gly
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5
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Leu Gly Cys Asn Ser Phe Arg Tyr
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<210> 474
<211> 17
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<400> 474
Cys Phe Gly Gly Arg Met Asp Arg Ile Gly Ala Gln Ser Gly Leu Gly
                5
Cys
<210> 475
<211> 22
<212> PRT
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<400> 475
Cys Phe Gly Gly Arg Met Asp Arg Ile Gly Ala Gln Ser Gly Leu Gly
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Cys Asn Ser Phe Arg Tyr
            20
<210> 476
<211> 28
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 476
   Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly
                                      10
   Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
               20
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   <210> 477
   <211> 11
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        Peptide
   <400> 477
   Arg Cys Gly Gly Arg Ile Asp Arg Ile Arg Cys
   <210> 478
   <211> 26
   <212> PRT
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        Peptide
   Arg Arg Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln
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Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
               20
                                   25
   <210> 479
   <211> 26
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   <223> Description of Artificial Sequence: Synthetic
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<400> 479
Arg Arg Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly Ala Gln
Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
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<210> 480
<211> 25
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Arg Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly Ala Gln Ser
                                    10
Gly Leu Gly Cys Asn Ser Phe Arg Tyr
            20
<210> 481
<211> 23
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     Peptide
<400> 481
Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly Ala Gln Ser Gly
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                                    10
                                                        15
Leu Gly Cys Asn Ser Phe Arg
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<210> 482
<211> 27
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 482
Ser Lys Ser Ser Ser Pro Cys Phe Gly Gly Lys Leu Asp Arg Ile Gly
                                   10
Ser Tyr Ser Gly Leu Gly Cys Asn Ser Arg Lys
           20
<210> 483
<211> 21
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 483
Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser Gly
                                    10
                                                         15
Leu Gly Cys Asn Ser
            20
<210> 484
<211> 23
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
   Peptide
<400> 484
Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser Gly
                                   10
Leu Gly Cys Asn Ser Phe Arg
            20
<210> 485
<211> 24
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<212> PRT
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<223> Description of Artificial Sequence: Synthetic
    Peptide
<400> 485
Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser Gly
1 5 10
Leu Gly Cys Asn Ser Phe Arg Tyr
          20
<210> 486
<211> 28
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    Peptide
<400> 486
Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly
 1 5
                                10
Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
               25
            20
<210> 487
<211> 24
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 487
Arg Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser
Gly Leu Gly Cys Asn Ser Phe Arg
            20
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<210> 488
<211> 25
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 488
Arg Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser
                  5
                                     10
                                                          15
Gly Leu Gly Cys Asn Ser Phe Arg Tyr
                                 25
             20
<210> 489
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<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 489
Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly
Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr Ser Leu Arg Arg
             20
Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly Ala Gln Ser Gly
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                             40
Leu Gly Cys Asn Ser Phe Arg Tyr
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<210> 490
<211> 32
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
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Peptide

<400> 490

Ala Gly Pro Arg Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Ile
1 5 10 15

Asp Arg Ile Gly Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
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<210> 491

<211> 27

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 491

Ser Lys Ser Ser Ser Pro Cys Phe Gly Gly Lys Leu Asp Arg Ile Gly
1 5 10 15

Ser Tyr Ser Gly Leu Gly Cys Asn Ser Arg Lys 20 25

<210> 492

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 492

Asn Pro Met Tyr Asn Ala Val Ser Asn Ala Asp Leu Met Asp Phe Lys

1 10 15

<210> 493

<211> 15

<212> PRT

<213> Artificial Sequence

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<220>
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        Peptide
  <400> 493
  Arg Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Cys
                   5
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                                                           15
  <210> 494
  <211> 21
  <212> PRT
  <213> Artificial Sequence
 <220>
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        Peptide
  <400> 494
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                                      10
 Asn Ser Phe Arg Tyr
              20
  <210> 495
  <211> 28
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Synthetic
        Peptide
<400> 495
  Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Met Asp Arg Ile Gly
                                      10
                                                           15
  Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
              20
  <210> 496
  <211> 21
  <212> PRT
  <213> Artificial Sequence
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<220>
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     Peptide
<400> 496
Phe Ala Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser Gly Leu Gly Cys
        5
                                  10
                                                     15
Asn Ser Phe Arg Tyr
           20
<210> 497
<211> 13
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 497
Ser Ser Asp Arg Ser Ala Leu Leu Lys Ser Lys Leu Arg
1 5
                      10
<210> 498
<211> 30
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 498
Asn Pro Met Tyr Asn Ala Val Ser Asn Ala Asp Leu Met Asp Phe Lys
                                 10
Asn Leu Leu Asp His Leu Glu Glu Lys Met Pro Leu Glu Asp
                    25
<210> 499
<211> 37
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Glu Val Val Pro Pro Gln Val Leu Ser Glu Pro Asn Glu Glu Ala Gly
                                    10
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Ala Ala Leu Ser Pro Leu Pro Glu Val Pro Pro Trp Thr Gly Glu Val
                                 25
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Ser Pro Ala Gln Arg
         35
 <210> 500
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       Peptide
  Ser Ser Asp Arg Ser Ala Leu Leu Lys Ser Lys Leu Arg Ala Leu Leu
                                      10
                    5
  Thr Ala Pro Arg
               20
   <210> 501
   <211> 27
   <212> PRT
   <213> Artificial Sequence
   <220>
   <223> Description of Artificial Sequence: Synthetic
         Peptide .
    Ser Lys Ser Ser Pro Cys Phe Gly Gly Lys Leu Asp Arg Ile Gly
                                    10
                      5
    Ser Tyr Ser Gly Leu Gly Cys Asn Ser Arg Lys
                                   25
                 20
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<211> 24
  <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 502
 Tyr Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser
                                       10
 Gly Leu Gly Cys Asn Ser Phe Arg
              20
 <210> 503
 <211> 21
 <212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 503
Tyr Ser Ser Asp Arg Ser Ala Leu Leu Lys Ser Lys Leu Arg Ala Leu
                                     10
                                                          15
Leu Thr Ala Pro Arg
             20
<210> 504
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 504
Thr Ala Pro Arg Ser Leu Arg Arg Ser Ser Cys Phe Gly Gly Arg Met
 1
                  5
                                     10
                                                         15
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<210> 502

Asp Arg Ile Gly Ala Gln Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr 20 25 30

<210> 505

<211> 24

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 505

Tyr Ser Ser Cys Phe Gly Gly Arg Ile Asp Arg Ile Gly Ala Gln Ser

1 5 10 15

Gly Leu Gly Cys Asn Ser Phe Arg

20

<210> 506

<211> 35

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 506

Ser Phe Asn Ser Cys Phe Gly Asn Arg Ile Glu Arg Ile Gly Ser Trp

1 5 10 15

Ser Gly Leu Gly Cys Asn Asn Val Lys Thr Gly Asn Lys Lys Arg Ile 20 25 30

Phe Gly Asn

35

<210> 507

<211> 32

<212> PRT

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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 507
Ser Pro Lys Met Met His Lys Ser Gly Cys Phe Gly Arg Arg Leu Asp
                                     10
Arg Ile Gly Ser Leu Ser Gly Leu Gly Cys Asn Val Leu Arg Lys Tyr
                                25
<210> 508
<211> 22
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 508
Gly Trp Asn Arg Gly Cys Phe Gly Leu Lys Leu Asp Arg Ile Gly Ser
                                    10
                                                         15
Leu Ser Gly Leu Gly Cys
            20
<210> 509
<211> 32
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 509
Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
                                     10
Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His
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20 25 30

<210> 510 <211> 45 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 510 Ser Gln Gly Ser Thr Leu Arg Val Gln Gln Arg Pro Gln Asn Ser Lys Val Thr His Ile Ser Ser Cys Phe Gly His Lys Ile Asp Arg Ile Gly 20 25 Ser Val Ser Arg Leu Gly Cys Asn Ala Leu Lys Leu Leu 35 40 <210> 511 <211> 26 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 511

Gly Leu Gly Cys Asn Val Leu Arg Arg Tyr
20 25

<210> 512 <211> 32 <212> PRT <213> Artificial Sequence

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Asp Ser Gly Cys Phe Gly Arg Arg Leu Asp Arg Ile Gly Ser Leu Ser

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<220>

<223> Description of Artificial Sequence: Synthetic Peptide <400> 512 Ser Pro Lys Thr Met Arg Asp Ser Gly Cys Phe Gly Arg Arg Leu Asp 10 Arg Ile Gly Ser Leu Ser Gly Leu Gly Cys Asn Val Leu Arg Arg Tyr 25 <210> 513 <211> 32 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 513 Asn Ser Lys Met Ala His Ser Ser Ser Cys Phe Gly Gln Lys Ile Asp 10 Arg Ile Gly Ala Val Ser Arg Leu Gly Cys Asp Gly Leu Arg Leu Phe 20 25 <210> 514 <211> 45 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide Ser Gln Asp Ser Ala Phe Arg Ile Gln Glu Arg Leu Arg Asn Ser Lys 5 10

Met Ala His Ser Ser Ser Cys Phe Gly Gln Lys Ile Asp Arg Ile Gly
20 25 30

Ala Val Ser Arg Leu Gly Cys Asp Gly Leu Arg Leu Phe 35 40 45

<210> 515

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<400> 515

Tyr Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met

1 5 10 15

Asp Arg Leu Ser Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg 20 25 30

His

<210> 516

<211> 32 .

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
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<400> 516

Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp 1 5 10 15

Arg Ile Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His
20 25 30

<210> 517

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<211> 22
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 517
Gly Leu Ser Lys Gly Cys Phe Gly Leu Lys Leu Asp Arg Ile Gly Ser
                 5
                                   10
Met Ser Gly Leu Gly Cys
            20
<210> 518
<211> 22
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      Peptide
Gly Leu Ser Arg Ser Cys Phe Gly Val Lys Leu Asp Arg Ile Gly Ser
                                   10
                                                        15
                 5
Met Ser Gly Leu Gly Cys
            20
<210> 519
<211> 53
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
Asp Leu Arg Val Asp Thr Lys Ser Arg Ala Ala Trp Ala Arg Leu Leu
                5
Gln Glu His Pro Asn Ala Arg Lys Tyr Lys Gly Ala Asn Lys Lys Gly
             20
                               25
                                                   30
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Leu Ser Lys Gly Cys Phe Gly Leu Lys Leu Asp Arg Ile Gly Ser Met
                           40
Ser Gly Leu Gly Cys
    50
<210> 520
<211> 53
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
Asp Leu Arg Val Asp Thr Lys Ser Arg Ala Ala Trp Ala Arg Leu Leu
                 5
                                                        15
His Glu His Pro Asn Ala Arg Lys Tyr Lys Gly Gly Asn Lys Lys Gly
                                25
            20
Leu Ser Lys Gly Cys Phe Gly Leu Lys Leu Asp Arg Ile Gly Ser Met
Ser Gly Leu Gly Cys
    50
<210> 521
<211> 27
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 521
Lys Pro Gly Thr Pro Pro Lys Val Pro Arg Thr Pro Pro Gly Glu Glu
                5
                                    10
Leu Ala Glu Pro Gln Ala Ala Gly Gly Asn Gln
             20
                                 25
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<210> 522
 <211> 21
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 <213> Artificial Sequence
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 522
 Gly Asp Lys Thr Pro Gly Gly Gly Gly Ala Asn Leu Lys Gly Asp Arg
                                      10
 Ser Arg Leu Leu Arg
             20′,
 <210> 523
 <211> 27
 <212> PRT
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 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 523
 Gly Leu Ser Lys Gly Cys Phe Gly Leu Lys Leu Asp Arg Ile Gly Ser
 Met Ser Gly Leu Gly Cys Asn Ser Phe Arg Tyr
             20
                                  25
 <210> 524
 <211> 23
<212> PRT
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       Peptide
 <400> 524
 Tyr Gly Leu Ser Lys Gly Cys Phe Gly Leu Lys Leu Asp Arg Ile Gly
                   5
                                     10
                                                          15
  1
 Ser Met Ser Gly Leu Gly Cys
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<210> 525
<211> 7
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
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Asp Ala Phe Val Ala Leu Met
1 5
<210> 526
<211> 7
<212> PRT
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     Peptide
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Asp Ala Phe Val Ala Leu Met
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<211> 11
<212> PRT
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<400> 527
Glu Pro Ser Lys Asp Ala Phe Ile Gly Leu Met
               5
                                   10
<210> 528
<211> 9
<212> PRT
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 528
Gly Pro Ser Gly Phe Tyr Gly Val Arg
<210> 529
<211> 10
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 529
Ala Pro Leu Ser Gly Phe Tyr Gly Val Arg
                5
<210> 530
<211> 7
<212> PRT
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      Peptide
<400> 530
Asp Ser Phe Val Gly Leu Met
1
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<210> 531
<211> 10
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 531
His Lys Thr Asp Ser Phe Val Gly Leu Met
1 5
<210> 532
<211> 10
<212> PRT
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     Peptide
<400> 532
His Lys Ile Asn Ser Phe Val Gly Leu Met
1 . 5
<210> 533
<211> 11
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 533
Tyr His Lys Thr Asp Ser Phe Val Gly Leu Met
1
          5
<210> 534
<211> 10
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 534
His Lys Thr Asp Ser Tyr Val Gly Leu Met
              5
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<210> 535

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<211> 7
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     Peptide
<400> 535
Lys Asp Ser Phe Val Gly Met
<210> 536
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     Peptide
<400> 536
Arg Ala Trp Phe Pro Pro
      5
<210> 537
<211> 6
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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<400> 537
Ala Ala Trp Phe Pro Pro
<210> 538
<211> 6
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Peptide
<400> 538
Ala Ala Trp Phe Pro Pro
<210> 539
<211> 7
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 539
Asp Ser Phe Val Ala Leu Met
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<210> 540
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 540
Asp Ser Phe Val Gly Leu
 1 5
<210> 541
<211> 7
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 541
Asp Ser Phe Trp Ala Leu Met
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<210> 542
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 542
Asp Tyr Trp Val Trp Trp Arg
    5
1
<210> 543
<211> 7
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 543
Asp Tyr Trp Val Trp Trp Lys
1 5
<210> 544
<211> 10
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 544
Asp Met His Asp Phe Phe Val Gly Leu Met
       5
<210> 545
<211> 9
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 545
Asp Met His Asp Phe Phe Gly Leu Met
      5
<210> 546
<211> 10
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<213> Artificial Sequence
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     Peptide
<400> 546
Asp Met His Asp Phe Phe Pro Gly Leu Met
                5
                                10
<210> 547
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 547
Tyr Asp Met His Asp Phe Phe Val Gly Leu Met
1
                 5
                                   10
<210> 548
<211> 9
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 548
Asp Pro His Asp Phe Trp Val Trp Leu
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Ile Arg Val His Pro Arg Gly Asn Leu Trp Ala Thr Gly His Phe Met

20 25 30

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<210> 552
<211> 5
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<213> Artificial Sequence
<220>
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      Peptide
<400> 552
Cys Tyr Trp Val Cys
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      Peptide
<400> 553
Gly Asn His Trp Ala Val Gly His Leu Met
                5
<210> 554
<211> 6
<212> PRT
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      Peptide
<400> 554
Lys Ile Pro Tyr Ile Leu
                  5
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<210> 555
<211> 8
<212> PRT
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<220>
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     Peptide
<400> 555
Tyr Phe Leu Phe Arg Pro Arg Asn
1 5
<210> 556
<211> 25
<212> PRT
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     Peptide
<400> 556
Phe Lys Val Asp Glu Glu Phe Gln Gly Pro Ile Val Ser Gln Asn Arg
                 5
                                    10
Arg Tyr Phe Leu Phe Arg Pro Arg Asn
            20
<210> 557
<211> 23
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 557
Tyr Lys Val Asn Glu Tyr Gln Gly Pro Val Ala Pro Ser Gly Gly Phe
                                    10
                                                        15
Phe Leu Phe Arg Pro Arg Asn
             20
```

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<210> 558
<211> 21
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 558
Asp Ala Gly His Gly Gln Ile Ser His Lys Arg His Lys Thr Asp Ser
                                    10
Phe Val Gly Leu Met
             20
<210> 559
<211> 10
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 559
Pro Asn Pro Asp Glu Phe Val Gly Leu Met
                 5
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<210> 560
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Gly Ser Phe Arg
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Arg Ser
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Leu Gly Ser Pro Ser Arg Ser
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Leu Ser Asn His Arg Gly Ser Leu Arg
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Leu Asp Ile Ile Trp
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Asp Ile Ile Trp
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Leu Asp Ile Ile Trp
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Leu Asp Ile Ile Trp
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Ala Ile Ile Trp
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Leu Asn Ile Ile Trp
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Gly Leu Gly Ser Pro Arg Ser
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<210> 717
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Leu Asp Ile Ile Trp
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Leu Asp Ile Ile Trp
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<210> 724

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Ser Pro Val Leu Trp
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Tyr Thr Asn Gly Leu Gln Val Asn Phe His Ser Pro Arg Ser Gly Gln
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Arg Cys Trp Ala Ala Arg Thr Gln Val Glu Lys Arg Leu
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Gln Asp Val Ile Trp
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1 5
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Tyr Pro Phe Pro Gly
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Tyr Pro Phe Pro Gly
1 5
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Tyr Phe
1
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1
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Tyr Phe Pro
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Tyr Phe Pro
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<210> 760
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1 .
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<210> 764
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1
<210> 765
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Tyr Phe Pro
1
<210> 766
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Gly Phe Leu Arg Arg Ile Arg Pro Lys Leu Lys Trp Asp Asn Gln

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 1 5
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                                    10
Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala His Lys Lys Gly Gln
            20
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                                    10
Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala His
            20
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                                    10
Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala Tyr Lys Lys Gly Glu
             20
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  Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala Tyr
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           5 10
     Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala His Lys Lys Gly Gln
                                   25
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Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala His
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              5
                                10
Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala His Lys Lys Gly Gln
                         25
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 1 5 10 15
Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala Tyr Lys Lys Gly Glu
                             25 .
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Tyr Lys Lys Gly Glu
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Tyr Gly Gly Phe Met Thr Ser Glu Lys Ser Gln Thr Pro Leu Val Thr
           5
Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala
            20 · 25
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                                   10
Leu Phe Lys Asn Ala Ile Ile Lys Asn Ala Tyr
                                25
            20
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Ile Ile Lys Asn Ala Tyr Lys Lys Gly Glu
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Leu Phe Lys Asn Ala Ile Val Lys Asn Ala His Lys Lys Gly Gln
             20
                                  25
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Leu Phe Lys Asn Ala Ile Ile Lys Asn Val His Lys Lys Gly Gln
                               25
           20
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           20 25
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<210> 843
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                                                         15
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<210> 861
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Tyr Gln Lys Arg Tyr Gly Gly Phe Leu

20 25

<210> 894

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

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Glu Glu Glu Ala Asn Gly Gly Glu Val Leu Gly Lys Arg Tyr Gly Gly 20 25 30

Phe Met

<210> 895

<211> 19

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Peptide

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Ser Ser Glu Val Ala Gly Glu Gly Asp Gly Asp Ser Met Gly His Glu 1 5 10 15

Asp Leu Tyr

<210> 896

<211> 6

<212> PRT

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<220>

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Tyr Pro Pro Pro
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Ala Ala Ala Tyr Gly Gly Phe Leu
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<210> 903
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Tyr Gln
<210> 905
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Tyr Pro Phe Pro Pro Leu
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   Peptide
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Tyr Val Pro Phe Pro Pro Phe
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Tyr Arg Phe Lys
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, <210> 909
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 Tyr Met Phe His Leu Met Asp
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 <210> 910
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<211> 7
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Tyr Ala Phe Glu Val Val Gly
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<210> 912
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Tyr Pro Trp Phe
 1
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Peptide
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Tyr Pro Trp Phe Phe
<210> 914
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Tyr
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<210> 916
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<210> 918
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<212> PRT
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Arg Phe
<210> 919
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Gln
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Tyr Pro Leu Gly
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<210> 922
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Tyr Pro Trp Gly
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<211> 9
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Pro Leu
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<210> 926
<211> 17
<212> PRT
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Phe Gly Gly Phe Thr Gly Ala Arg Lys Ser Ala Arg Lys Leu Ala Asn
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15

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Gln
<210> 927
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Tyr Gly Gly Phe Met Arg Arg Val
<210> 928
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Ala Lys Ser Gln Gly Gly Ser Asn
<210> 929
<211> 8
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<400> 929
Leu Glu Asp Gly Pro Lys Phe Leu
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1

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<210> 930
<211> 5
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Arg Lys Asp Val Tyr
<210> 931
<211> 13
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Gly Glu Gln Arg Lys Asp Val Tyr Val Gln Leu Tyr Leu
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<210> 932
<211> 28
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Lys Glu Lys Lys Glu Val Val Glu Glu Ala Glu Asn
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<210> 933
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                 5
 <210> 934
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 Lys Glu Lys Lys Glu Val Val Glu Glu Ala Glu Asn
              20
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 Arg Phe Gly Thr Cys Thr Val Gln Lys Leu Ala His Gln Ile Tyr Gln
              20
                                 25
 Phe Thr Asp Lys Asp Lys Asp Asn Val Ala Pro Arg Ser Lys Ile Ser
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                              40
                                                  45
 Pro Gln Gly Tyr
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<210> 937
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Gln Ile Tyr Phe Thr Asp Lys Asp Asn Val Ala Pro Arg Ser Lys Ile
                               25
Ser Pro Gln Gly Tyr
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<210> 938
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Thr Val Gln Lys Leu Ala His Gln Ile Tyr Gln Phe Thr Asp Lys Asp
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1 5 10 15

Lys Asp Asn Val Ala Pro Arg Ser Lys Ile Ser Pro Gln Gly Tyr \$20\$ \$25\$ 30

<210> 939

<211> 48

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 939

Glu Leu Arg Met Ser Ser Ser Tyr Pro Thr Gly Leu Ala Asp Val Lys

1 5 10 15

Ala Gly Pro Ala Gln Thr Leu Ile Arg Pro Gln Asp Met Lys Gly Ala 20 25 30

Ser Arg Ser Pro Glu Asp Ser Ser Pro Asp Ala Ala Arg Ile Arg Val\$35\$ 40 45

<210> 940

<211> 33

<212> PRT

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<400> 940

Ser Leu Pro Glu Ala Gly Pro Gly Arg Thr Leu Val Ser Ser Lys Pro 1 5 10 15

Gln Ala His Gly Ala Pro Ala Pro Pro Ser Gly Ser Ala Pro His Phe 20 25 30

Leu

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<210> 941
<211> 52
<212> PRT
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Phe Gly Thr Cys Thr Val Gln Lys Leu Ala His Gln Ile Tyr Gln Phe
            20
                                25
Thr Asp Lys Asp Gly Val Ala Pro Arg Ser Lys Ile Ser Lys Ile Ser
                             40
                                                45
Pro Gln Gly Tyr
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Ala Leu Ser Arg
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Peptide

<400> 943

Tyr Arg Gln Ser Met Asn Gln Gly Ser Arg Ser Thr Gly Cys Arg Phe
1 5 10 15

Gly Thr Cys Thr Met Gln Lys Leu Ala His Gln Ile Tyr Gln Ile Tyr
20 25 30

Gln Phe Thr Asp Lys Asp Lys Asp Gly Met Ala Pro Arg Asn Lys Ile 35 40 45

Ser Pro Gln Gly Tyr 50

<210> 944

<211> 40

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 944

Ser Thr Gly Cys Arg Phe Gly Thr Cys Thr Met Gln Lys Leu Ala His 1 5 10 15

Gln Ile Tyr Gln Phe Thr Asp Lys Asp Lys Asp Gly Met Ala Pro Arg \$20\$ \$25\$ 30

Asn Lys Ile Ser Pro Gln Gly Tyr 35 40

<210> 945

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Peptide

<400> 945

Ala Arg Leu Asp Thr Ser Ser Gln Phe Arg Lys Lys Trp Asn Lys Trp

1 10 15

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Ala Leu Ser Arg
<210> 946
<211> 13
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<400> 946
Ala Pro Ser Gly Ala Gln Arg Leu Tyr Gly Phe Gly Leu
 1 5
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<210> 948
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<212> PRT
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     Peptide
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Gly Gly Ser Leu Tyr Ser Phe Gly Leu
 1 5 .
<210> 949
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315

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<211> 8
<212> PRT
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      Peptide
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Asp Arg Leu Tyr Ser Phe Gly Leu
<210> 950
<211> 17
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Val His His Gln Lys Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn
                                    10
                                                         15
                 5
Lys
<210> 951
<211> 11
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<400> 951
Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu Met
        5
<210> 952
<211> 5
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 952
Arg Glu Arg Met Ser
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<210> 953
<211> 17
<212> PRT
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<400> 953
Ala Lys Glu Arg Leu Glu Ala Lys His Arg Glu Arg Met Ser Gln Val
                                   10
Met
<210> 954
<211> 43
<212> PRT
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     Peptide
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Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys
                                   10
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Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
                         25
Gly Leu Met Val Gly Gly Val Val Ile Ala Thr
        35
                            40
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<210> 955

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<211> 42
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     Peptide
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Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys
                5
                                   10
Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
            20
                                25
Gly Leu Met Val Gly Gly Val Val Ile Ala
                           40
<210> 956
<211> 40
<212> PRT
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     Peptide
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                 5
                                    10
Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
Gly Leu Met Val Gly Gly Val Val
 35 40
<210> 957
<211> 11
<212> PRT
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      Peptide
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Tyr Glu Val His His Gln Lys Leu Val Phe Phe
<210> 958
<211> 14
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 958
Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu Met
                                    10
<210> 959
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<400> 959
Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys
                  5
                                     10
Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys
             20
                                 25
<210> 960
<211> 40
<212> PRT
<213> Artificial Sequence
<220>
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      Peptide
<400> 960
Val Val Gly Gly Val Met Leu Gly Ile Ile Ala Gly Lys Asn Ser Gly
 1
                  5
                                     10
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Val Asp Glu Ala Phe Phe Val Leu Lys Gln His His Val Glu Tyr Gly 25 20 Ser Asp His Arg Phe Glu Ala Asp 35 <210> 961 <211> 42 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 961 Asp Ala Glu Phe Gly His Asp Ser Gly Phe Glu Val Arg His Gln Lys 5 Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val Ile Ala 35 40 <210> 962 <211> 40 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide Asp Ala Glu Phe Gly His Asp Ser Gly Phe Glu Val Arg His Gln Lys 1 5 10 Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile 25 Gly Leu Met Val Gly Gly Val Val

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Tyr Glu Val His His Gln Lys Leu Val Phe Phe
<210> 964
<211> 14
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Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu Met
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<211> 28
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Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys
<210> 966
<211> 40
<212> PRT
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<213> Artificial Sequence

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Val Asp Glu Ala Phe Phe Val Leu Lys Gln His His Val Glu Tyr Gly
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Ser Asp His Arg Phe Glu Ala Asp
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Gly Leu Met Val Gly Gly Val Val Ile Ala
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<210> 968
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<212> PRT
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<400> 968
Asp Ala Glu Phe Gly His Asp Ser Gly Phe Glu Val Arg His Gln Lys
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Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
             20
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Gly Leu Met Val Gly Gly Val Val
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<210> 969
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Ile Gly Leu Met
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<210> 975

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<211> 38
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 Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys
 Leu Val Phe Phe Ala Glu Asp Val Gly Ser ..sn Lys Gly Ala Ile Ile
                                 25
 Gly Leu Met Val Gly Gly
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  <210> 976
  <211> 28
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  1 5
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              20
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 Val Ala Gln Lys Thr Val Glu Gly Ala Gly Ser Ile Ala Ala Ala Thr
             20
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 Gly Phe Val
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 <210> 981
 <211> 24
 <212> PRT
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 Gly Leu Met Val Gly Gly Val Val
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 <210> 982
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  <210> 983
  <211> 17
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Lys
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Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu Met
 1 5 10
<210> 985
<211> 5
<212> PRT
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Arg Glu Arg Met Ser
<210> 986
<211> 17
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Met
<210> 987
<211> 43
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Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys
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Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
                                 25
Gly Leu Met Val Gly Gly Val Val Ile Ala Thr
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                                     10
Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
             20
                                 25
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Gly Leu Met Val Gly Gly Val Val Ile Ala
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Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
                               25
            20
Gly Leu Met Val Gly Gly Val Val
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<210> 990
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Tyr Glu Val His His Gln Lys Leu Val Phe Phe
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Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu Met
 1 5
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                                  10
Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys
            20
                               25
<210> 993
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                                   10
Val Asp Glu Ala Phe Phe Val Leu Lys Gln His His V\epsilon Glu Tyr Gly
                        25
            20
Ser Asp His Arg Phe Glu Ala Asp
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<210> 994
<211> 42
<212> PRT
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Peptide

<400> 994

Asp Ala Glu Phe Gly His Asp Ser Gly Phe Glu Val Arg His Gln Lys

1 10 15

Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile 20 25 30

Gly Leu Met Val Gly Gly Val Val Ile Ala 35 40

<210> 995

<211> 40

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 995

Asp Ala Glu Phe Gly His Asp Ser Gly Phe Glu Val Arg His Gln Lys

1 5 10 15

Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile 20 25 30

Gly Leu Met Val Gly Gly Val Val
35 40

<210> 996

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 996

Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu
1 5 10

<210> 997

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<211> 5
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Ile Ile Gly Leu Met
<210> 998
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Ile Gly Leu Met
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<210> 999
<211> 11
<212> PRT
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<400> 999
Met Leu Gly Ile Ile Ala Gly Lys Asn Ser Gly
<210> 1000
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334

<210> 1003 <211> 28 <212> PRT

<213> Artificial Sequence

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<400> 1003
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Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys
            20
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Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Gln Val His His Gln Lys
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                      10
<210> 1005
<211> 40
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1005
Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val His His Gln Lys
Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
                              25
Gly Leu Met Val Gly Gly Val Val
        35
                            40
<210> 1006
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335

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<211> 35
<212> PRT
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<220>
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     Peptide
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His Asp Ser Gly Tyr Glu Val His His Gln Lys Leu Val Phe Phe Ala
                                   10
Gln Asp Val Gly Ser Asn Lys Gly Ala Ile Ile Gly Leu Met Val Gly
                                25
Gly Val Val
    . 35
<210> 1007
<211> 35
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     Peptide
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Glu Gln Val Thr Asn Val Gly Gly Ala Val Val Thr Gly Val Thr Ala
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Val Ala Gln Lys Thr Val Glu Gly Ala Gly Ser Ile Ala Ala Ala Thr
                                25
Gly Phe Val
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<210> 1008
<211> 24
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1008
Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys Gly Ala Ile Ile
     5
                                  10
                                                      15
Gly Leu Met Val Gly Gly Val Val
            20
<210> 1009
<211> 11
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Gly Tyr Val Ile Ile Lys Pro Leu Val Trp Val
<210> 1010
<211> 63
<212> PRT
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     Peptide
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Arg Leu Glu Lys Arg Asp Val Lys Ala Val Cys Ser Gln Glu Ala Met
                                  10
Thr Gly Pro Cys Arg Ala Val Met Pro Arg Trp Tyr Phe Asp Leu Ser
20
                       25
Lys Gly Lys Cys Val Arg Phe Ile Tyr Gly Gly Cys Gly Gly Asn Arg
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Asn Asn Phe Glu Ser Glu Asp Tyr Cys Met Ala Val Cys Lys Ala
     50
                        55
<210> 1011
<211> 39
<212> PRT
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<213> Artificial Sequence

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Asn Ser Ser Cys Gly Glu Gly Asn His Leu Pro Thr Thr Pro Cys Tyr
                                     10
                5
Leu Gln Trp Gly Thr His Arg Glu Phe Leu Arg Arg Phe Ser Ile Trp
                                 25
Asn His Gly His Leu His Met
        35
<210> 1012
<211> 39
<212> PRT
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      Peptide
<400> 1012
Ala Gly Arg Gly Lys Gln Gly Gly Lys Val Arg Ala Lys Ala Lys Thr
                                    10
Arg Ser Ser Arg Ala Gly Leu Gln Phe Pro Val Gly Arg Val His Arg
                                25
             20
Leu Leu Arg Lys Gly Asn Tyr
        35
<210> 1013
<211> 21
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1013
Thr Arg Ser Ser Arg Ala Gly Leu Gln Phe Pro Val Gly Arg Val His
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1 5 10 15

Arg Leu Leu Arg Lys 20

<210> 1014

<211> 37

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1014

Lys Trp Lys Leu Phe Lys Lys Ile Glu Lys Val Gly Gln Asn Ile Arg

1 5 10 15

Asp Gly Ile Ile Lys Ala Gly Pro Ala Val Ala Val Val Gly Gln Ala 20 25 30

Thr Gln Ile Ala Lys 35

<210> 1015

<211> 32

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1015

Ser Trp Leu Ser Lys Thr Ala Lys Lys Leu Glu Asn Ser Ala Lys Lys

1 5 10 15

Arg Ile Ser Glu Gly Ile Ala Ile Ala Ile Gln Gly Gly Pro Arg Cys 20 25 30

<210> 1016

<211> 31

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     Peptide
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Gly Ile Met Ser Ile Val Lys Asp Val Ala Lys Asn Ala Ala Lys Gly
                                 10
1 5
Ala Leu Ser Thr Leu Ser Thr Leu Ser Cys Lys Leu Ala Lys Thr
                             25
           20
<210> 1017
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Phe Leu Gly Ala Leu Phe Lys Val Ala Ser Lys Val Leu Pro Ser Val
                      10
       5
Lys Cys Ala Ile Thr Lys Lys Cys
           20
<210> 1018
<211> 13
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Ile Leu Pro Trp Lys Trp Pro Trp Pro Trp Arg Arg
<210> 1019
<211> 18
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340

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<212> PRT
<213> Artificial Sequence
<220>
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      Peptide
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Arg Gly Gly Arg Leu Cys Tyr Cys Arg Arg Arg Phe Cys Val Cys Val
                  5
                                    10
Gly Arg
<210> 1020
<211> 18
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Arg Arg Trp Cys Tyr Arg Lys Cys Tyr Lys Gly Tyr Cys Tyr Arg Lys
                                   10
Cys Arg
<210> 1021
<211> 35
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      Peptide
<400> 1021
Arg Trp Lys Ile Phe Lys Lys Ile Glu Lys Val Gly Gln Asn Ile Arg
                                    10
Asp Gly Ile Val Lys Ala Gly Pro Ala Val Ala Val Val Gly Gln Ala
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Ala Thr Ile
       35
<210> 1022
<211> 12
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Arg Leu Cys Arg Ile Val Val Ile Arg Val Cys Arg
        5
                       10
<210> 1023
<211> 21
<212> PRT
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Thr Arg Ser Ser Arg Ala Gly Leu Gln Phe Pro Val Gly Arg Val His
               5
                                  10
                                                     15
Arg Leu Leu Arg Lys
            20
<210> 1024
<211> 14
<212> PRT
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     Peptide
<400> 1024
Tyr Pro Pro Gly Pro Leu Ala Pro Pro Gln Pro Phe Gly Pro
                5
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<210> 1025
<211> 22
<212> PRT
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Val Asn Pro Gly Val Val Val Arg Ile Ser Gln Lys Gly Leu Asp Tyr
     5
                               10
Ala Ser Gln Gln Gly Arg
           20
<210> 1026
<211> 15
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Ala Asp Val Leu Thr Thr Gly Ala Gly Asn Pro Val Gly Asp Lys
1 5 10 15
<210> 1027
<211> 42
<212> PRT
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     Peptide
<400> 1027
Met Ala Ala Val Ser Met Ser Val Ala Leu Arg Gln Ala Leu Trp Gly
                      10
Arg Arg Val Ala Thr Val Ala Ala Val Ser Val Ser Lys Val Ser Thr
                         25 30
           20
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Arg Ser Leu Ser Thr Ser Thr Trp Arg Leu
       35 40
<210> 1028
<211> 23
<212> PRT
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     Peptide
<400> 1028
Gly Ile Gly Lys Phe Lys His Ser Ala Gly Lys Phe Gly Lys Ala Phe
                                  10
Val Gly Glu Ile Met Lys Ser
            20
<210> 1029
<211> 29
<212> PRT
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     Peptide
<400> 1029
Leu Arg Asp Leu Val Ser Tyr Cys Arg Ala Arg Gly Lys Gly Arg Glu
                5
                                   10
Arg Met Asn Gly Thr Arg Lys Gly His Leu Leu Tyr Met
            20
                                25
<210> 1030
<211> 4
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      Peptide
<400> 1030
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Ser Gln Asn Tyr
   1
 <210> 1031
 <211> 12
 <212> PRT
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       Peptide
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 Arg Leu Cys Arg Ile Val Val Ile Arg Val Cys Arg
                 5
 <210> 1032
 <211> 25
 <212> PRT
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       Peptide
 <400> 1032
 Asn Gln Gly Arg His Phe Cys Gly Gly Ala Leu Ile His Ala Arg Phe
   1
                                     10
                                                          15
 Val Met Thr Ala Ala Ser Cys Phe Gln
              20
 <210> 1033
<211> 2
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        Peptide
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<211> 18
<212> PRT
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<400> 1034
Cys Asn Leu Ala Val Ala Ala Ser His Ile Tyr Gln Asn Gln Phe
Val Gln
<210> 1035
<211> 19
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1035
Tyr Arg Val Arg Phe Leu Ala Lys Glu Asn Val Thr Gln Asp Ala Glu
                                   10
                                                      15
                5
Asp Asn Cys
<210> 1036
<211> 15
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
Lys Trp Lys Leu Phe Lys Lys Ile Gly Ala Val Leu Lys Val Leu
        5
                                    10
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<210> 1034

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<210> 1037
          <211> 35
          <212> PRT
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                Peptide
          <400> 1037
          Lys Trp Lys Val Phe Lys Lys Ile Glu Lys Met Gly Arg Asn Ile Arg
                5
          Asn Gly Ile Val Lys Ala Gly Pro Ala Ile Ala Val Leu Gly Glu Ala
                      20
          Lys Ala Leu
          <210> 1038
          <211> 12
          <212> PRT
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          <220>
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                Peptide
          <400> 1038
          Thr Tyr Ile Cys Glu Val Glu Asp Gln Lys Glu Glu
                         5
<211> 29
         <210> 1039
          <212> PRT
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               Peptide
          <400> 1039
          Cys Tyr Cys Arg Ile Pro Ala Cys Ile Ala Gly Glu Arg Arg Tyr Gly
                          5
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Thr Cys Ile Tyr Gln Gly Arg Leu Trp Ala Phe Cys Cys 20 <210> 1040 <211> 34 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1040 Ala Leu Trp Lys Thr Met Leu Lys Lys Leu Gly Thr Met Ala Leu His 10 Ala Gly Lys Ala Ala Leu Gly Ala Ala Asp Thr Ile Ser Gln Gly 25 30 20 Thr Gln <210> 1041 <211> 24 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1041 Gly Val Leu Ser Asn Val Ile Gly Tyr Leu Lys Lys Leu Gly Thr Gly 5 10 Ala Leu Asn Ala Val Leu Lys Gln 20 <210> 1042 <211> 6 <212> PRT <213> Artificial Sequence <220>

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Val Glu Pro Ile Pro Tyr
<210> 1043
<211> 11
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Glu Arg Glu Val Arg
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<400> 1045

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<210> 1046
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Val Gly Glu Ile Met Lys Ser
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Gly Ile Gly Lys Phe Leu His Ser Ala Lys Lys Phe Gly Lys Ala Phe
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Val Gly \mathbb{G}^{\eta}u Ile Met Asn Ser
             20
<210> 1048
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Tyr Leu Val Leu Phe Phe Tyr Pro Leu Asp Phe Thr Phe Val Cys Pro
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Thr Glu Ile Ile Cys Pro Thr Glu Ile Ile Gly
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<210> 1049
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Leu Val Gln Ala Phe Gln Gly Lys Val Asn Val Phe Leu Gln Phe
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<210> 1050
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Gly Gly Lys
<210> 1051
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Asp Glu Val Asp
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Asp Glu Val Asp
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<210> 1064
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Phe Tyr Phe Ala Ser
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<210> 1071
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Ala Asn Arg Leu Phe Gly Glu Lys
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Val Asn Lys
<210> 1073
<211> 14
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Gly Asn Thr Ala Ala Gln Met Ala Gln Ile Leu Ser Phe Asn
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                                    10
<210> 1074
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Gln Gly Leu Trp Leu Met Asn Val Leu Arg Val Gly Trp His His His
Leu Gln Pro Arg Thr Val Pro Glu Asn
          20
<210> 1075
<211> 35
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1 5 10 15
Leu Ile Met His Lys Ile Thr Asn Cys Ile Leu Phe Phe Gly Arg Phe
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Ser Ser Pro
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<210> 1076
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Ala Pro Arg Leu Arg Phe Tyr Ser Leu
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Ala Pro Arg Leu Arg Phe Tyr
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Arg Leu Arg Phe His
<210> 1080
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Arg Leu Arg Phe Asp
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Glu Glu Arg Leu His Ser Met
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<210> 1082
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Gln Lys Leu Gly Asn Gln Trp Ala Val Gly His Leu Met
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<210> 1084

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<211> 14
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Glu Gln Arg Leu Gly Asn Gln Trp Ala Val Gly His Leu Met
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<210> 1085
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                 5
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Phe Gln Trp Ala Val Gly His Leu
<210> 1087
<211> 13
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Met Pro Leu Pro Pro His Pro Gly His Pro Gly Tyr Ile Asn Phe Ser
                                     10
Tyr Glu Val Leu Thr
             20
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Pro Phe Tyr Gly Pro Val
<210> 1093
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<400> 1093
Tyr Leu Tyr Gln Trp Leu Gly Ala Pro Val Pro Tyr Pro Asp Pro Leu
                                    10
Glu Pro Arg Arg Val Cys Leu Asn Pro Asp Cys Asp Glu Leu Ala Asp
             20
                                 25
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His Ile Gly Phe Gln Glu Ala Tyr Arg Arg Phe Tyr Gly Pro Val
        35
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Leu Val Val Tyr Pro Trp
<210> 1095
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Tyr Leu Tyr Gln Trp Leu Gly Ala Pro Val Pro Tyr Pro Asp Pro Leu
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               5 10
                                                     15
Pro Arg Arg Val Cys Leu Asn Pro Asp Cys Asp Glu Leu Ala Asp His
            20
                              25
Ile Gly Phe Gln Glu Ala Tyr Arg Arg Phe Tyr Gly Pro Val
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<210> 1096
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Gly Phe Gln Glu Ala Tyr Arg Arg Phe Tyr Gly Pro Val
 1 5 10
<210> 1097
<211> 13
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Pro Tyr Gln Glu Ala Phe Arg Arg Phe Phe Gly Pro Val
<210> 1098
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Val Pro Ile Tyr Glu Lys Lys Tyr Gly Gln Val Pro Met Cys Asp Ala
 1
                5 10
                                                    15
Gly Glu Gln Cys Ala Val Arg Lys Gly Ala Arg Ile Gly Lys Leu Cys
            20
                             25
Asp Cys Pro Arg Gly Thr Ser Cys Asn Ser Phe Leu Leu Lys Cys Leu
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<210> 1099
<211> 48
<212> PRT
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Peptide

<400> 1099

Val Pro Ile Tyr Glu Lys Lys Tyr Gly Gln Val Pro Met Cys Asp Ala 1 5 10 15

Gly Glu Gln Cys Ala Val Arg Lys Gly Ala Arg Ile Gly Lys Leu Cys 20 25 30

Asp Cys Pro Arg Gly Thr Ser Cys Asn Ser Phe Leu Leu Lys Cys Leu 35 40 45

<210> 1100

<211> 48

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Val Pro Ile Tyr Glu Lys Lys Tyr Gly Gln Val Pro Met Cys Asp Ala 1 5 10 15

Gly Glu Gln Cys Ala Val Arg Lys Gly Ala Arg Ile Gly Lys Leu Cys 20 25 30

Asp Cys Pro Arg Gly Thr Ser Cys Asn Ser Phe Leu Leu Lys Cys Leu 35 40 45

<210> 1101

<211> 28

<212> PRT

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<220>

<223> Description of Artificial Sequence: Synthetic Peptide

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Ala Gly Ala Thr Val Gln Val Thr Leu Asp Gly Val Pro Ala Ala Ala
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                  5
Pro Gly Gln Pro Ala Gln Leu Gln Leu Asn Ala Thr
            20
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Phe Leu Phe Leu Phe
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<210> 1103
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Ala Cys Tyr Cys Arg Ile Pro Ala Cys Ile Ala Gly Glu Arg Arg Tyr
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Gly Thr Cys Ile Tyr Gln Gly Arg Leu Trp Ala Phe Cys Cys
                               25
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<210> 1104
<211> 3
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      Peptide
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Met Leu Phe
  1
<210> 1105
<211> 4
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<210> 1106
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Met Leu Phe Lys
 1
<210> 1107
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Trp His Val Ala Ala Asn
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<210> 1108

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<211> 25
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Met Leu Thr Glu Leu Glu Lys Ala Leu Asn Ser Ile Ile Asp Val Tyr
                 5
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His Lys Tyr Ser Leu Ile Lys Gly Asn
             20
<210> 1109
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His Trp Asp Thr Thr Gln Ser Leu Lys Gln Leu Glu Glu Arg Ala Ala
                                    10
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Trp Asn Val
<210> 1110
<211> 20
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      Peptide
Gly Pro Val Ser Ala Val Leu Thr Glu Leu Arg Cys Thr Cys Leu Arg
                 5
                                    10
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Val Thr Leu Arg
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370

20

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 Ile Gln Arg Thr Pro Lys Ile Gln Val Tyr Ser Arg His Pro Ala Glu
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 Asn Gly Lys Ser Asn
             20
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                 5
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  Tyr Ser Asp Asp Glu Gly Ala Ser Trp Ser Asp Leu Asp Ile Val Ser
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 Phe Ser Lys
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Val Leu
<210> 1115
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<210> 1116
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Cys Asn Thr Gly Gln Leu Cys Pro Val Glu
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<210> 1117
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372

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Leu Glu
<210> 1118
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Leu His Glu Trp Thr Lys Pro Glu Asn Leu Asp Phe Ile Glu Val Asn
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                                  10
Val Leu Pro
<210> 1119
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Val Leu Glu Leu Pro Tyr Gln Gly Glu Glu Leu Ser Met Val Ile Leu
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               5
Leu Pro Asp Asp Ile Glu Asp Glu Ser Thr Gly Leu Lys
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            20
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<211> 19
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Thr Tyr Gly Ala Asp Leu Ala Ser Val Asp Phe Gln His Ala Ser Glu
                                    10
                  5
Asp Ala Arg
<210> 1121
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Glu Arg Pro Pro Leu Gln Gln Pro Pro His Arg Asp
                  5
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<210> 1122
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                  5
                                     10
Cys Lys Asn Phe Phe Trp Lys Thr Phe Ser Ser Cys Lys
             20
                                  25
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                                    10
Asp Ala Thr Gly
            20
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                                     10
Asp Ser Ser Ser Thr Ala Phe Glu Gly Gly Thr Pro Glu Leu Ser
             20
                                 25
<210> 1125
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<210> 1126
<211> 4
<212> PRT
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Arg Gly Asp Ser
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<210> 1127
<211> 4
<212> PRT
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<400> 1127
Arg Gly Asp Val
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<210> 1128
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<400> 1128
Arg Gly Glu Ser
 1
<210> 1129
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<210> 1130
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        5
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<210> 1132
<211> 49
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      Peptide
<400> 1132
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378

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  <210> 1137
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<210> 1158
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<210> 1162

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  <210> 1168
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   <210> 1174
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                                                            15
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Val Val Lys Leu Lys Asp Glu Asn Ser Gln Leu Lys Ser Glu Val Ser
                20
                                    25
   Lys Leu Arg Cys Gln Leu Ala Lys Lys
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<210> 1184
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Ser Asp Arg Asn Phe Leu Arg Phe
1 5
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Asp Pro Phe Leu Arg Phe
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Glu Ile Leu Glu Val Pro Ser Thr
                 5
<210> 1188
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Gly Trp Thr Leu Asn Ser Ala Gly Tyr Leu Leu Gly Pro His Ala Val
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Gly Asn His Arg Ser Phe Ser Asp Lys Asn Gly Leu Thr Ser
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Gly Asn His
<210> 1190
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                                     10
Ala Leu Ser Ala Ser Ala Gly Leu Trp Ser Pro Ala Lys Glu
                                 25
             20
<210> 1191
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Glu Leu Arg Pro Glu Asp Asp Met Lys Pro Gly Ser Phe Asp Arg Ser
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Ile Pro Glu Asn Asn Ile Met Arg
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Thr Ile Ile Glu Phe Leu Ser Phe Leu His Leu Lys Glu Ala Gly Ala
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Leu Asp Arg Leu Leu Asp Leu Pro Ala Ala Ala Ser Ser Glu Asp Ile
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Glu Arg Ser
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Asp Asn His Arg Ser Phe His Asp Lys Tyr Gly Leu Ala
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<210> 1194
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 <210> 1195
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                                           15
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 Asp Asn His Arg Ser Phe Ser Asp Lys His Gly Leu Thr
                              25
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 <211> 40
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1 5
 Gly Pro His Ala Ile Asp Asn His Arg Ser Phe Ser Asp Lys His Gly
            20
                           25
 Leu Thr Gly Lys Arg Glu Leu Pro
                           40
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Phe Ser Pro Phe Arg
            20
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Leu Ala Leu Ala
            20
<210> 1199
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Pro Gln Gln Trp Phe Trp Leu Leu
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                           10
   Gln Ser Glu Asp Lys Ala Ile Arg Thr Ile Met Glu Phe Leu Ala Phe
               20
                                  25
   Leu His Leu Lys Glu Ala Gly Ala Leu
                              40
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   <211> 26
   <212> PRT
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           5
                          10
   Phe Leu His Leu Lys Glu Ala Gly Ala Leu
<210> 1202
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   <213> Artificial Sequence
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1 5 10

<210> 1209

<211> 120

<212> PRT

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<220>

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<400> 1209

Lys Lys Lys Asp Lys Val Lys Lys Gly Gly Pro Gly Ser Glu Cys Ala 1 5 10 15

Glu Trp Ala Trp Gly Pro Cys Thr Pro Ser Ser Lys Asp Cys Gly Val 20 25 30

Gly Phe Arg Glu Gly Thr Cys Gly Ala Gln Thr Gln Arg Ile Arg Cys
35 40 45

Arg Val Pro Cys Asn Trp Lys Lys Glu Phe Gly Ala Asp Cys Lys 50 55 60

Phe Glu Asn Trp Gly Ala Cys Asp Gly Gly Thr Gly Thr Lys Val Arg 65 70 75 80

Gln Gly Thr Leu Lys Lys Ala Arg Tyr Asn Ala Gln Cys Gln Glu Thr 85 90 95

Ile Arg Val Thr Lys Pro Cys Thr Pro Lys Thr Lys Ala Lys Ala Lys 100 105 110

Ala Lys Lys Gly Lys Gly Lys Asp 115 120

<210> 1210

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<220>

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Peptide

<400> 1210

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Gly Thr Lys Val Arg Gln Gly Thr Leu Lys Lys Ala Arg Tyr Asn Ala
                   25
Gln Cys Gln Glu Thr Ile Arg Val Thr Lys Pro Cys Thr Pro Lys Thr
        35
                            40
Lys Ala Lys Ala Lys Lys Gly Lys Gly Lys Asp
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Cys His Ser Gly Tyr Val Gly Val Arg Cys
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<210> 1212
<211> 13
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Ala Asn Phe Leu Val Trp Glu Ile Val Arg Lys Lys Pro
                5
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<210> 1213
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Peptide

<400> 1213

Val Val Ser His Phe Asn Asp Cys Pro Asp Ser His Thr Gln Phe Cys

1 10 15

Phe His Gly Thr Cys Arg Phe Leu Val Gln Glu Asp Lys Pro Ala Cys
20 25 30

Val Cys His Ser Gly Tyr Val Gly Ala Arg Cys Glu His Ala Asp Leu 35 40 45

Leu Ala 50

<210> 1214

<211> 50

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1214

Val Val Ser His Phe Asn Lys Cys Pro Asp Ser His Thr Gln Tyr Cys

1 5 10 15

Phe His Gly Thr Cys Arg Phe Leu Val Gln Glu Glu Lys Pro Ala Cys
20 25 30

Val Cys His Ser Gly Tyr Val Gly Val Arg Cys Glu His Ala Asp Leu 35 40 45

Leu Ala 50

<210> 1215

<211> 10

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<223> Description of Artificial Sequence: Synthetic Peptide

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Cys His Ser Gly Tyr Val Gly Val Arg Cys
                5
<210> 1216
<211> 10
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Cys His Ser Gly Tyr Val Gly Val Arg Cys
1 5
<210> 1217
<211> 6
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Pro Pro Gly His Phe Lys
 1
                5
<210> 1218
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Arg Thr Gly Gln Tyr Lys
<210> 1219
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<211> 11
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Phe Asn Leu Pro Leu Gly Asn Tyr Lys Lys Pro
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<210> 1220
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                                10
                                                        15
Phe Lys Asp Pro Lys Arg Leu Tyr
            20
<210> 1221
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<210> 1222
<211> 12
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                                10
<210> 1223
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Asp Val Val Asp Ala Asp Glu Tyr Leu Ile Pro Gln
                5
<210> 1224
<211> 12
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    Peptide
<400> 1224
Cys Tyr Ala Ala Pro Leu Lys Pro Ala Lys Ser Cys
         5
<210> 1225
<211> 18
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1225
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Tyr Phe Asn Lys Pro Thr Gly Tyr Gly Ser Ser Ser Arg Arg Ala Pro
1 5
                     10
Gln Thr
<210> 1226
<211> 14
<212> PRT
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     Peptide
<400> 1226
Ala Leu Leu Glu Thr Tyr Cys Ala Thr Pro Ala Lys Ser Glu
       5
                                 10
<210> 1227
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     Peptide
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Gly Tyr Gly Ser Ser Ser Arg Arg Ala Pro Gln Thr
                                 10
               5
<210> 1228
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Ser Arg Val Ser Arg Arg Ser Arg
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<210> 1229
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<223> Description of Artificial Sequence: Synthetic
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<400> 1229
Tyr Ser Arg Val Ser Arg Arg Ser Arg
<210> 1230
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<210> 1231
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Ser Val Arg Val Glu Gln Val Val Lys Pro Pro Gln Asp Lys Thr Glu
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Ser Glu Asn Thr Ser Asp
             20
<210> 1232
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Glu Trp Gln Trp Ser Val Cys Val
            20
<210> 1233
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Ser Ala Gln Thr Asn Arg His Ile Leu Arg Phe Asn Arg Pro Phe
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<210> 1235
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<212> PRT
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     Peptide
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Ile Pro Val Lys Gln Ala Val His Gly Gln Phe Leu Leu Pro Lys Gln
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Glu Lys
<210> 1236
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1236
Leu Ala Gly Glu Thr Gly Gln Glu Ala Ala Pro Leu Asp Gly Val Leu
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Ala Asn
<210> 1237
<211> 14
<212> PRT
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 Peptide
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Ala Leu Lys Arg Gln Gly Arg Thr Leu Tyr Gly Phe Gly Gly
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                5
<210> 1238
<211> 30
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<220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1238 Gly Met Asp Val Leu Gly Arg Pro Lys Ile Pro Leu Glu Thr Pro Ala 10 Tyr Thr Gly Gln Pro Trp His Cys Gln His Cys Phe Leu Leu <210> 1239 <211> 43 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1239 Met Asn Thr Ile Thr Ile Cys Lys Phe Asp Val Leu Asp Ala Glu Leu 10 Leu Ser Thr Val Glu Gly Gly Tyr Ser Gly Lys Asp Cys Leu Lys Asp 25 Met Gly Gly Tyr Ala Leu Ala Gly Ala Gly Ser <210> 1240 <211> 38 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1240 Gly Ala Asp Lys Thr Val Lys Gly Pro Asp Gly Leu Thr Ala Leu Glu 10 Ala Thr Asp Asn Gln Ala Ile Asp Tyr Gly Gly Phe Met Glu Val Val 20 25

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Tyr Val Asp Ala Thr Lys
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<210> 1241
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<212> PRT
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Cys Lys Gln Leu Gln Arg Asp Arg Gln Val Tyr Arg Ala Thr His Arg
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<210> 1242
<211> 17
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Cys Glu Gly Asn Val Arg Val Ser Arg Glu Leu Ala Gly His Thr Gly
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Tyr
<210> 1243
<211> 16
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      Peptide
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Cys Gly Ala Gly Glu Ser Gly Lys Ser Thr Ile Val Lys Gln Met Lys
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                                                         15
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Cys Asn Leu Lys Glu Asp Gly Ile Ser Ala Ala Lys Asp Val Lys
                                   10
<210> 1245
<211> 16
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1245
Cys Lys Gln Leu Gln Lys Asp Lys Gln Val Tyr Arg Ala Thr His Arg
                          10
<210> 1246
<211> 11
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Glu Glu Gln Gly Met Leu Pro Glu Asp Leu Ser
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<210> 1247
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Pro Gly Thr Cys Glu Ile Cys Ala Tyr Ala Ala Cys Thr Gly Cys
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<210> 1248
<211> 15
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<400> 1248
Pro Asn Thr Cys Glu Ile Cys Ala Tyr Ala Ala Cys Thr Gly Cys
               5
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<210> 1249
<211> 16
<212> PRT
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     Peptide
<400> 1249
Asn Asp Asp Cys Glu Leu Cys Val Asn Val Ala Cys Thr Gly Cys Leu
               5
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                                                       15
              <210> 1250
<211> 32
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1250
Ser Thr Pro Leu Met Ser Trp Pro Trp Ser Pro Ser Ala Leu Arg Leu
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1 5 10 15

Leu Gln Arg Pro Pro Glu Glu Pro Ala Ala His Ala Asn Cys His Arg 20 25 30

<210> 1251

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1251

Tyr Ser Thr Pro Leu Met Ser Trp Pro Trp Ser Pro Ser Ala Leu Arg
1 5 10 15

Leu Leu Gln Arg Pro Pro Glu Glu Pro Ala Ala His Ala Asn Cys His
20 25 30

Arg

<210> 1252

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1252

His Asn Lys Gln Glu Gly Arg Asp His Asp Lys Ser Lys Gly His Phe

1 5 10 15

His Arg Val Val Ile His His Lys Gly Gly Lys Ala His Arg Gly
20 25 30

<210> 1253

<211> 32

<212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1253 Tyr His Asn Lys Gln Glu Gly Arg Asp His Asp Lys Ser Lys Gly His 5 Phe His Arg Val Val Ile His His Lys Gly Gly Lys Ala His Arg Gly 20 25 <210> 1254 <211> 32 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1254 Ser Thr Ala Pro Leu Pro Trp Pro Trp Ser Pro Ala Ala Leu Arg Leu 5 10 Leu Gln Arg Pro Pro Glu Glu Pro Ala Val His Ala Asp Cys His Arg 20 25 <210> 1255 <211> 33 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1255

Tyr Ser Thr Ala Pro Leu Pro Trp Pro Trp Ser Pro Ala Ala Leu Arg 10 Leu Leu Gln Arg Pro Pro Glu Glu Pro Ala Val His Ala Asp Cys His 20 25 Arg <210> 1256 <211> 17 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1256 Val Gln Gly Glu Thr Ser Asn Asp Lys Ile Pro Val Ala Leu Gly Leu 10 Lys <210> 1257 <211> 72 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1257 Ala Pro Val Ala Asn Glu Leu Arg Cys Gln Cys Leu Gln Thr Val Ala 10 Gly Ile His Phe Lys Asn Ile Gln Ser Leu Lys Val Met Pro Pro Gly 20 Pro His Cys Thr Gln Thr Glu Val Ile Ala Thr Leu Lys Asn Gly Arg 35 40 45 Glu Ala Cys Leu Asp Pro Glu Ala Pro Met Val Gln Lys Ile Val Gln

60

55

50

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Lys Met Leu Lys Gly Val Pro Lys
65
                 70
<210> 1258
<211> 9
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1258
Val Gln Gly Glu Glu Ser Asn Asp Lys
<210> 1259
<211> 13
<212> PRT
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     Peptide
<400> 1259
Ile Leu Asn Gly Ile Asn Asn Tyr Lys Asn Pro Lys Leu
                5
                                  10
<210> 1260
<211> 22
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1260
Phe Val Gln Gly Glu Ala Ile Pro Met Ser Ile Pro Pro Glu Asp Lys
                 5
                       10
                                                      15
Ile Pro Val Ala Leu Gly
            20
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<210> 1261
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1261
Ala Pro Val Pro Pro Gly Glu Asp Ser Lys Asp Val Ala
      5
<210> 1262
<211> 21
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1262
Met Val Lys Gln Ile Glu Ser Lys Thr Ala Phe Gln Glu Ala Leu Asp
               5
                                  10
Ala Ala Gly Asp Lys
           20
<210> 1263
<211> 10
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1263
Ala Tyr Val His Asp Ala Pro Val Arg Ser
 1 5 10
<210> 1264
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<211> 7
<212> PRT
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<400> 1264
Pro Arg Lys Leu Tyr Asp Lys
           5
<210> 1265
<211> 12
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      Peptide
<400> 1265
Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr Lys
<210> 1266
<211> 24
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1266
Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Ala Tyr Thr Thr Asn
                                    10
                5
Pro Arg Lys Leu Tyr Asp Tyr Lys
             20
<210> 1267
<211> 12
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1267
Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Lys
                5
<210> 1268
<211> 8
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1268
Pro Arg Lys Leu Tyr Asp Tyr Lys
      5
<210> 1269
<211> 12
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1269
Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr Lys
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              5
                                    10
<210> 1270
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1270
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Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr
 1 5
<210> 1271
<211> 24
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1271
Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Ala Tyr Thr Thr Asn
                                 10
Pro Arg Lys Leu Tyr Asp Tyr Lys
            20
<210> 1272
<211> 23
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1272
Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Ala Tyr Thr Thr Asn
         5
                                  10
Pro Arg Lys Leu Tyr Asp Tyr
           20
<210> 1273
<211> 7
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1273
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Arg Lys Leu Tyr Asp Tyr Lys
1 5
<210> 1274
<211> 6
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1274
Arg Lys Leu Tyr Asp Tyr
<210> 1275
<211> 7
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1275
Pro Arg Lys Leu Tyr Asp Lys
                5
<210> 1276
<211> 6
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1276
Pro Arg Lys Leu Tyr Asp
<210> 1277
<211> 8
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<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1277
Pro Arg Lys Leu Tyr Asp Tyr Lys
      5
<210> 1278
<211> 4
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     Peptide
<400> 1278
Asp Gly Glu Ala
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<210> 1279
<211> 6
<212> PRT
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      Peptide
<400> 1279
Leu Gly Thr Ile Pro Gly
                5 .
<210> 1280
<211> 23
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1280
Asn Ile Ser Ser Glu Glu Lys Ala Ser Trp Thr Arg Pro Glu Lys Gln
                                    10
Glu Thr Leu Asp Gly His Met
             20
<210> 1281
<211> 17
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1281
Phe Asn Leu Asp Val Arg Phe Leu Val Val Lys Glu Ala Val Asn Pro
                  5
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Gly
<210> 1282
<211> 17
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1282
Ala Pro Arg Gln Arg Gln Thr Leu Val Leu Phe Pro Gly Asp Leu Arg
            . 5
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Thr
<210> 1283
<211> 24
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1283
Cys Ser Cys Ser Pro Val His Pro Gln Gln Ala Phe Cys Asn Ala Asp
Val Val Ile Arg Ala Lys Ala Val
            20
<210> 1284
<211> 24
<212> PRT
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     Peptide
<400> 1284
Cys Ser Cys Ser Pro Val His Pro Gln Gln Ala Phe Cys Asn Ala Asp
                                   10
Val Val Ile Arg Ala Lys Ala Val
            20
<210> 1285
<211> 13
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1285
Asn Val Ile Gln Ile Ser Asn Asp Leu Glu Asn Leu Arg
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                5
                                   10
<210> 1286
<211> 35
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
Val Pro Ile Gln Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys Thr
       5
                                   10
Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr Gln Ser Val Ser Ser
                                25
Lys Gln Lys
<210> 1287
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1287
Tyr Lys Val Gln Asp Asp Thr Lys Thr Leu Ile Lys Thr Ile Val
                 5
                                   10
<210> 1288
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
 Peptide
<400> 1288
Ser Cys Ser Leu Pro Gln Thr Ser Gly Leu Gln Lys Pro Glu Ser
               5
                                    10
<210> 1289
<211> 10
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1289
Glu Asp Val Asp His Val Phe Leu Arg Phe
         5
<210> 1290
<211> 7
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1290
Thr Ser Phe Thr Pro Arg Leu
                5
<210> 1291
<211> 8
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1291
Asp Pro Ala Phe Asn Ser Trp Gly
<210> 1292
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1292
Asp Pro Gly Phe Ser Ser Trp Gly
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<210> 1293
<211> 8
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1293
Asp Gln Gly Phe Asn Ser Trp Gly
<210> 1294
<211> 8
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1294
Asp Ala Ser Phe His Ser Trp Gly
<210> 1295
<211> 8
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1295
Gly Ser Gly Phe Ser Ser Trp Gly
        5
<210> 1296
<211> 8
<212> PRT
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1296
Asp Pro Ala Phe Ser Ser Trp Gly
1
        5
<210> 1297
<211> 8
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1297
Gly Ala Ser Phe Tyr Ser Trp Gly
 1
                5
<210> 1298
<211> 10
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1298
Ile Ile Gly Gly Arg Glu Ser Arg Pro His
                     10
       5
<210> 1299
<211> 38
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1299
His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Gln
                                10
Met Ala Val Lys Lys Tyr Leu Ala Ala Val Leu Gly Lys Arg Tyr Lys
                             25
Gln Arg Val Lys Asn Lys
       35
<210> 1300
<211> 30
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1300
His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Arg Gln
                                 10
               5
Leu Ala Val Arg Arg Tyr Leu Ala Ala Val Leu Gly Lys Arg
                              25
<210> 1301
<211> 21
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
    Peptide
                    Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Met Ala Val Lys Lys Tyr
                      10
Leu Ala Ala Val Leu
            20
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432

<210> 1302 <211> 38 <212> PRT

<213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1302 His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Gln 5 10 Met Ala Val Lys Lys Tyr Leu Ala Ala Val Leu Gly Lys Arg Tyr Lys 25 Gln Arg Ile Lys Asn Lys 35 <210> 1303 <211> 27 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1303 His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Gln 10 Met Ala Val Lys Lys Tyr Leu Ala Ala Val Leu 20 <210> 1304 <211> 22 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1304 Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Gln Met Ala Val Lys Lys 5 10

433

Tyr Leu Ala Ala Val Leu

20

<210> 1305 <211> 38 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1305 His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Gln 10 5 Met Ala Val Lys Lys Tyr Leu Ala Ala Val Leu Gly Lys Arg Tyr Lys 20 25 Gln Arg Val Lys Asn Lys 35 <210> 1306 <211> 33 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1306 Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Gln Met Ala Val Lys Lys 15 5 Tyr Leu Ala Ala Val Leu Gly Lys Arg Tyr Lys Gln Arg Val Lys Asn 25 Lys

<210> 1307

<211> 23

<212> PRT

<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1307
Gln Met Ala Val Lys Lys Tyr Leu Ala Ala Val Leu Gly Lys Arg Tyr
                                   10
Lys Gln Arg Val Lys Asn Lys
            20
<210> 1308
<211> 11
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1308
Gly Lys Arg Tyr Lys Gln Arg Val Lys Asn Lys
<210> 1309
<211> 8
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1309
Tyr Lys Gln Arg Val Lys Asn Lys
        . 5
<210> 1310
<211> 29
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1310
Asp Val Ala His Gly Ile Leu Asn Glu Ala Tyr Arg Lys Val Leu Asp
                  5
                                     10
Gln Leu Ser Ala Gly Lys His Leu Gln Ser Leu Val Ala
<210> 1311
<211> 29
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1311
Asp Val Ala His Glu Ile Leu Asn Glu Ala Tyr Arg Lys Val Leu Asp
                                     10
Gln Leu Ser Ala Arg Lys Tyr Leu Gln Ser Met Val Ala
                                 25
            20
<210> 1312
<211> 20
<212> PRT
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      Peptide
<400> 1312
Ser Asp Glu Asp Ser Asp Gly Asp Arg Pro Gln Ala Ser Pro Gly Leu
                  5
                                    10
                                                         15
Gly Pro Gly Pro
             20
<210> 1313
<211> 52
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1313
Gly Glu Ser Arg Ser Glu Ala Leu Ala Val Asp Gly Ala Gly Lys Pro
Gly Ala Glu Glu Ala Gln Asp Pro Glu Gly Lys Gly Glu Gln Glu His
            20
                              25
Ser Gln Gln Lys Glu Glu Glu Glu Met Ala Val Val Pro Gln Gly
        35
                          40
Leu Phe Arg Gly
    50
<210> 1314
<211> 29
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1314
Pro Glu Gly Lys Gly Glu Gln Glu His Ser Gln Gln Lys Glu Glu Glu
               5
                                      15
Glu Glu Met Ala Val Val Pro Gln Gly Leu Phe Arg Gly
           20
                              25
<210> 1315
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1315
Glu Glu Glu Glu Met Ala Val Pro Gln Gly Leu Phe Arg Gly
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<210> 1316
<211> 49
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1316
Gly Trp Pro Gln Ala Pro Ala Met Asp Gly Ala Gly Lys Thr Gly Ala
                                    10
Glu Glu Ala Gln Pro Pro Glu Gly Lys Gly Ala Arg Glu His Ser Arg
             20
                                 25
Gln Glu Glu Glu Glu Thr Ala Gly Ala Pro Gln Gly Leu Phe Arg
                             40
                                                 45
Gly
<210> 1317
<211> 48
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1317
Gly Trp Pro Gln Ala Pro Ala Asp Gly Ala Gly Lys Thr Gly Ala Glu
                                     10
Glu Ala Gln Pro Pro Glu Gly Lys Gly Ala Arg Glu His Ser Arg Gln
                                 25
Glu Glu Glu Glu Glu Thr Ala Gly Ala Pro Gln Gly Leu Phe Arg Gly
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<210> 1318

40

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<211> 49
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1318
Tyr Gly Trp Pro Gln Ala Pro Ala Asp Gly Ala Gly Lys Thr Gly Ala
                                   10
Glu Glu Ala Gln Pro Pro Glu Gly Lys Gly Ala Arg Glu His Ser Arg
                                25
Gln Glu Glu Glu Glu Thr Ala Gly Ala Pro Gln Gly Leu Phe Arg
                            40
Gly
<210> 1319
<211> 50
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1319
Tyr Gly Trp Pro Gln Ala Pro Ala Met Asp Gly Ala Gly Lys Thr Gly
                                   10
Ala Glu Glu Ala Gln Pro Pro Glu Gly Lys Gly Ala Arg Glu His Ser
                 . 25
Arg Gln Glu Glu Glu Glu Thr Ala Gly Ala Pro Gln Gly Leu Phe
         35
                            40
                                               45
Arg Gly
     50
<210> 1320
<211> 19
<212> PRT
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439

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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1320
Leu Ser Phe Arg Ala Pro Ala Tyr Gly Phe Arg Gly Pro Gly Leu Gln
                                   10
        5
Leu Arg Arg
<210> 1321
<211> 51
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1321
Asp Asp Gly Gln Ser Glu Ser Gln Ala Val Asn Gly Lys Thr Gly Ala
                                    10
Ser Glu Ala Val Pro Ser Glu Gly Lys Gly Glu Leu Glu His Ser Gln
                                25
Gln Glu Glu Asp Gly Glu Glu Ala Met Ala Gly Pro Pro Gln Gly Leu
                            40
Phe Pro Gly
     50
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Met Pro Pro Arg Arg Asp Ala Met Pro
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Thr Met Leu Ala
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447

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Lys Asn
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1 5
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Glu Met Leu Ala
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Arg Gln Val Pro Leu Ser Ala Arg Gly
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Ala Asp Leu Asn
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Pro Ser Thr Gly
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Leu Leu Pro Arg Gly Gly Leu Ala Ala Leu
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Arg His Asn Pro Thr Gly Thr Val Leu Phe Met Gly Gln Val Met Glu
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Pro
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Gly Thr Asp Cys
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Ser Ser Ala Ser Cys Cys Gln Leu

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<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Peptide

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Thr Lys Cys Cys Arg Gly Arg Pro Cys Arg Cys Ser Met Ile Gly Thr
20 25 30

Asn Cys Glu Cys Thr Pro Arg Leu Ile Met Glu Gly Leu Ser Phe Ala 35 40 45

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<212> PRT

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<220>

<223> Description of Artificial Sequence: Synthetic Peptide

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Arg Cys Cys Ser Gly Leu Lys Cys Lys Glu Leu Ser Ile Trp Asp Ser 20 25 30

Arg Cys Leu

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Gln His
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Lys Lys Gln Phe Ser Ser Phe Tyr Phe Lys Trp Thr Ala Lys Lys Cys
                . 25
Leu Pro Phe Leu Phe Ser Gly Cys Gly Gly Asn Ala Asn Arg Phe Gln
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                                       45
Thr Ile Gly Glu Cys Arg Lys Lys Cys Leu Gly Lys
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     5 10
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Cys
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 1 5
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              5
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Ser Gly Ser Cys Gly Arg Arg Gly Lys Cys
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Lys Ile Cys Gly Lys Asn
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                                             15
Cys Lys Ala Gln Phe Gly Gln Ser Ala Gly Ala Lys Cys Met Asn Gly
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Lys Cys Lys Cys Tyr Pro His
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                                    10
Cys Asp Asp Cys Tyr Thr Cys Arg Cys Gly Thr Pro Trp Gly Ala Asn
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Cys Arg Cys Asp Tyr Tyr Lys Ala Arg Cys Asp Thr
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Leu Leu Gly Lys Cys Ile Gly Asp Lys Cys Glu Cys Val Lys His
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<400> 1415
Ile Asn Trp Lys Gly Ile Ala Ala Met Ala Lys Lys Leu Leu
     5
<210> 1416
<211> 21
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
Phe Met Ser Ser His Gln Ser Gln Ala Ser Leu Glu Leu Ala Ile Lys
                       10
Gln Trp Gly Ser Gln
           20
<210> 1417
<211> 20
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1417
Phe Asn Lys Ile Thr Pro Asn Leu Ala Glu Phe Ala Phe Ser Leu Tyr
                         10
Arg Gln Leu Ala
            20
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<210> 1418
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1418
Trp Ser Pro Lys Glu Glu Asp Arg Ile Ile Pro Gly Gly Ile Tyr Asn
Ala Asp Leu Asn Asp Glu Trp Val Gln Arg Ala Leu His Phe Ala Ile
                                 25
             20
Ser Glu
<210> 1419
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1419
Tyr Thr Ser Leu Ile His Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln
Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
                                 25
             20
Trp Asn Trp Phe
         35
<210> 1420
<211> 38
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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Peptide

<400> 1420

Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu
1 5 10 15

Thr Val Trp Gln Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu 20 25 30

Arg Tyr Leu Lys Asp Gln 35

<210> 1421

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1421

Tyr Thr Asn Thr Ile Tyr Thr Leu Leu Glu Glu Ser Gln Asn Gln Gln 1 5 10 15

Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu 20 25 30

Trp Asn Trp Phe

<210> 1422

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1422

Tyr Thr Gly Ile Ile Tyr Asn Leu Leu Glu Glu Ser Gln Asn Gln Gln 1 5 10 15

Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Asn Leu 20 25 30

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Trp Asn Trp Phe
      35
<210> 1423
<211> 36
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
    Peptide
<400> 1423
Tyr Thr Ser Leu Ile Tyr Ser Leu Leu Glu Lys Ser Gln Thr Gln Gln
1 5
                    10
Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
                                   30
                           25
           20
Trp Asn Trp Phe
       35
<210> 1424
<211> 36
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
    Peptide
<400> 1424
Leu Glu Ala Asn Ile Ser Lys Ser Leu Glu Gln Ala Gln Ile Gln Gln
1 5 10 15
Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu Asn Ser Trp Asp Ile Phe
                          25
                                  30
           20
Gly Asn Trp Phe
      35
<210> 1425
<211> 36
<212> PRT
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475

<213> Artificial Sequence

<220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1425 Leu Glu Ala Asn Ile Ser Gln Ser Leu Glu Gln Ala Gln Ile Gln Gln 10 Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu Asn Ser Trp Asp Val Phe 25 Thr Asn Trp Leu 35 <210> 1426 <211> 41 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1426 Cys Gly Gly Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu 10 Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu 25 Ala Val Glu Arg Tyr Leu Lys Asp Gln <210> 1427 <211> 38 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1427 Gln Gln Leu Leu Asp Val Val Lys Arg Gln Gln Glu Met Leu Arg Leu

10 15 1 Thr Val Trp Gly Thr Lys Asn Leu Gln Ala Arg Val Thr Ala Ile Glu 20 25 Lys Tyr Leu Lys Asp Gln 35 <210> 1428 <211> 46 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1428 Tyr Thr Ser Val Ile Thr Ile Glu Leu Ser Asn Ile Lys Glu Asn Lys 10 Cys Asn Gly Ala Lys Val Lys Leu Ile Lys Gln Glu Leu Asp Lys Tyr 25 Lys Asn Ala Val Thr Glu Leu Gln Leu Leu Met Gln Ser Thr 40 <210> 1429 <211> 54 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1429 Ala Ser Gly Val Ala Val Ser Lys Val Leu His Leu Glu Gly Glu Val 10 Asn Lys Ile Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly Val Ser Val Leu Thr Ser Lys Val Leu Asp Leu Lys Asn Tyr

40

Ile Asp Lys Gln Leu Leu 50 <210> 1430 <211> 53 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1430 Gly Glu Pro Ile Ile Asn Phe Tyr Asp Pro Leu Val Phe Pro Ser Asp 5 . 10 Glu Phe Asp Ala Ser Ile Ser Gln Val Asn Glu Lys Ile Asn Gln Ser 25 20 Leu Ala Phe Ile Arg Lys Ser Asp Glu Leu Leu His Asn Val Asn Ala 40 Gly Lys Ser Thr Thr 50 <210> 1431 <211> 48 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1431 Tyr Thr Ser Val Ile Thr Ile Glu Leu Ser Asn Ile Lys Glu Asn Lys 1 ' 5 10 Cys Asn Gly Thr Asp Ala Lys Val Lys Leu Ile Lys Gln Glu Leu Asp 25 20 Lys Tyr Lys Asn Ala Val Thr Glu Leu Gln Leu Leu Met Gln Ser Thr 40 45 35

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<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1432
Tyr Thr Ser Val Ile Thr Ile Glu Leu Ser Asn Ile Lys Glu Asn Lys
Cys Asn Gly Asp Ala Lys Val Lys Leu Ile Lys Gln Glu Leu Asp Lys
                              25
Tyr Lys
<210> 1433
<211> 34
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1433
Thr Ser Val Ile Thr Ile Glu Leu Ser Asn Ile Lys Glu Asn Lys Cys
                                  10
Asn Gly Asp Ala Lys Val Lys Leu Ile Lys Gln Glu Leu Asp Lys Tyr
                25 30
Lys Asn
<210> 1434
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
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<210> 1432

<223> Description of Artificial Sequence: Synthetic

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Peptide
<400> 1434
Val Ile Thr Ile Glu Leu Ser Asn Ile Lys Glu Asn Lys Cys Asn Gly
                                     10
Asp Ala Lys Val Lys Leu Ile Lys Gln Glu Leu Asp Lys Tyr Lys Asn
             20
                                 25
Ala Val
<210> 1435
<211> 34
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1435
Val Ile Thr Ile Glu Leu Ser Asn Ile Lys Glu Asn Lys Met Asn Gly
Asp Ala Lys Val Lys Leu Ile Lys Gln Glu Leu Asp Lys Tyr Lys Asn
                                 25
             20
Ala Val
<210> 1436
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1436
Val Ala Val Ser Lys Val Leu His Leu Glu Gly Glu Val Asn Lys Ile
Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly Val
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20 25 30

Ser

<210> 1437

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1437

Ala Val Ser Lys Val Leu His Leu Glu Gly Glu Val Asn Lys Ile Ala 1 5 10 15

Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly Val Ser
20 25 30

Val

<210> 1438

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<400> 1438

Val Ser Lys Val Leu His Leu Glu Gly Glu Val Asn Lys Ile Ala Leu 1 5 10 15

Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly Val Ser Val $20 \hspace{1.5cm} 25 \hspace{1.5cm} 30 \hspace{1.5cm}$

Leu

<210> 1439

<211> 33

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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1439
Ser Lys Val Leu His Leu Glu Gly Glu Val Asn Lys Ile Ala Leu Leu
 1
                5
                                   10
                                                       15
Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly Val Ser Val Leu
            20
                                25
Thr
<210> 1440
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1440
Lys Val Leu His Leu Glu Gly Glu Val Asn Lys Ile Ala Leu Leu Ser
                5
                                    10
Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly Val Ser Val Leu Thr
            20
                                25
                                                    30
Ser
<210> 1441
<211> 33
<212> PRT
<213> Artificial Sequence '
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
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482

<400> 1441

Leu Glu Gly Glu Val Asn Lys Ile Ala Leu Leu Ser Thr Asn Lys Ala 10 Val Val Ser Leu Ser Asn Gly Val Ser Val Leu Thr Ser Lys Val Leu Asp <210> 1442 <211> 33 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1442 Gly Glu Val Asn Lys Ile Ala Leu Leu Ser Thr Asn Lys Ala Val Val 1 5 Ser Leu Ser Asn Gly Val Ser Val Leu Thr Ser Lys Val Leu Asp Leu 25 Lys <210> 1443 <211> 33 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1443 Glu Val Asn Lys Ile Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser 5 15 Leu Ser Asn Gly Val Ser Val Leu Thr Ser Lys Val Leu Asp Leu Lys

Asn

20

25

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<210> 1444
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1444
Val Asn Lys Ile Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu
                          . 10
Ser Asn Gly Val Ser Val Leu Thr Ser Lys Val Leu Asp Leu Lys Asn
                               25
Tyr
<210> 1445
<211> 33
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1445
Asn Lys Ile Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser
                                   10
Asn Gly Val Ser Val Leu Thr Ser Lys Val Leu Asp Leu Lys Asn Tyr
  20
                               25
Ile
<210> 1446
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1446
Lys Ile Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn
                                    10
Gly Val Ser Val Leu Thr Ser Lys Val Leu Asp Leu Lys Asn Tyr Ile
             20
                                 25
Asp
<210> 1447
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1447
Ile Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly
                                     10
Val Ser Val Leu Thr Ser Lys Val Leu Asp Leu Lys Asn Tyr Ile Asp
                                 25
Lys
<210> 1448
<211> 33
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1448
Ala Leu Leu Ser Thr Asn Lys Ala Val Val Ser Leu Ser Asn Gly Val
                                    10
Ser Val Leu Thr Ser Lys Val Leu Asp Leu Lys Asn Tyr Ile Asp Lys
```

20 25 30

Gln

<210> 1449

<211> 70

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1449

Gly Thr Ile Ala Leu Gly Val Ala Thr Ser Ala Gln Ile Thr Ala Ala 1 5 10 15

Val Ala Leu Val Glu Ala Lys Gln Ala Arg Ser Asp Ile Glu Lys Leu 20 25 30

Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln Ser 40 45

Ser Ile Gly Asn Leu Ile Val Ala Ile Lys Ser Val Gln Asp Tyr Val 50 55 60

Asn Lys Glu Ile Val Pro 65 70

<210> 1450

<211> 56

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1450

Tyr Thr Pro Asn Asp Ile Thr Leu Asn Asn Ser Val Ala Leu Asp Pro 1 5 10 15

Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu 20 25 30

Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile Gly 40 Asn Trp His Gln Ser Ser Thr Thr 50 55 <210> 1451 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1451 Thr Leu Asn Asn Ser Val Ala Leu Asp Pro Ile Asp Ile Ser Ile Glu 10 Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg 25 Arg Ser Asn 35 <210> 1452 <211> 35 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1452 Leu Asn Asn Ser Val Ala Leu Asp Pro Ile Asp Ile Ser Ile Glu Leu 10 Asn Lys Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg 20 25 Ser Asn Gln 35

<210> 1453

<211> 35

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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1453
Asn Asn Ser Val Ala Leu Asp Pro Ile Asp Ile Ser Ile Glu Leu Asn
                                   10
Lys Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser
            20
                               25
                                                   30
Asn Gln Lys
   35
<210> 1454
<211> 35
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1454
Asn Ser Val Ala Leu Asp Pro Ile Asp Ile Ser Ile Glu Leu Asn Lys
                                   10
Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn
                               25
Gln Lys Leu
         35
<210> 1455
<211> 35
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1455 Ser Val Ala Leu Asp Pro Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala 10 Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln 25 Lys Leu Asp <210> 1456 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1456 Val Ala Leu Asp Pro Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys 5 10 Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys 20 25 30 Leu Asp Ser 35 <210> 1457 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1457 Ala Leu Asp Pro Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser 5 Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu 25 30 20

Asp Ser Ile

35

<210> 1458 <211> 35 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1458 Leu Asp Pro Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp 5 10 Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp 25 20 Ser Ile Gly 35 <210> 1459 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide Asp Pro Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu 5 10 Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser 20 Ile Gly Asn 35 <210> 1460 <211> 35 <212> PRT

<213> Artificial Sequence

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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1460
Pro Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu
                      10
Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile
                           25
            20
Gly Asn Trp
       35
<210> 1461
<211> 35
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1461
Ile Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu
               5 10
                                                   15
Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile Gly
                              25
Asn Trp His
        35
<210> 1462
<211> 35
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1462
Asp Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu Ser
               5
                                 10
                                             15
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Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile Gly Asn 25 Trp His Gln 35 <210> 1463 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1463 Ile Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile Gly Asn Trp 25 His Gln Ser 35 <210> 1464 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1464 Ser Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile Gly Asn Trp His 25 20 Gln Ser Ser 35

<210> 1465

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<211> 35
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1465
Ile Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp
                                    10
                 5
Ile Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile Gly Asn Trp His Gln
                                25
Ser Ser Thr
       35
<210> 1466
<211> 35
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1466
Glu Leu Asn Lys Ala Lys Ser Asp Leu Glu Glu Ser Lys Glu Trp Ile
                5
                                    10
Arg Arg Ser Asn Gln Lys Leu Asp Ser Ile Gly Asn Trp His Gln Ser
                                25
Ser Thr Thr
        35
<210> 1467
<211> 35
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1467

Thr Ala Ala Val Ala Leu Val Glu Ala Lys Gln Ala Arg Ser Asp Ile 10 Glu Lys Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser. 25 Val Gln Ser 35 <210> 1468 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1468 Ala Val Ala Leu Val Glu Ala Lys Gln Ala Arg Ser Asp Ile Glu Lys 5 10 Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln 25 30 20 Ser Ser Ile 35 <210> 1469 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1469 Leu Val Glu Ala Lys Gln Ala Arg Ser Asp Ile Glu Lys Leu Lys Glu 5 10 Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile 20 25 30 Gly Asn Leu

35

<211> 35 <212> PRT

<213> Artificial Sequence

<210> 1470 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1470 Val Glu Ala Lys Gln Ala Arg Ser Asp Ile Glu Lys Leu Lys Glu Ala 5 10 Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile Gly 25 Asn Leu Ile 35 <210> 1471 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide Glu Ala Lys Gln Ala Arg Ser Asp Ile Glu Lys Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile Gly Asn 20 25 Leu Ile Val 35 <210> 1472

<220>

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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1472
Ala Lys Gln Ala Arg Ser Asp Ile Glu Lys Leu Lys Glu Ala Ile Arg
Asp Thr Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile Gly Asn Leu
                    25
Ile Val Ala
       35
<210> 1473
<211> 35
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1473
Lys Gln Ala Arg Ser Asp Ile Glu Lys Leu Lys Glu Ala Ile Arg Asp
                                  10
Thr Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile Gly Asn Leu Ile
            20
                               25
Val Ala Ile
         35
<210> 1474
<211> 35
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1474
Gln Ala Arg Ser Asp Ile Glu Lys Leu Lys Glu Ala Ile Arg Asp Thr
                5
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Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile Gly Asn Leu Ile Val 20 25 Ala Ile Lys 35 <210> 1475 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1475 Ala Arg Ser Asp Ile Glu Lys Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile Gly Asn Leu Ile Val Ala 25 Ile Lys Ser 35 <210> 1476 <211> 35 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1476 Arg Ser Asp Ile Glu Lys Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys 10 Ala Val Gln Ser Val Gln Ser Ser Ile Gly Asn Leu Ile Val Ala Ile Lys Ser Val 35

<210> 1477

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<211> 35
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1477
Ser Asp Ile Glu Lys Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala
                                    10
Val Gln Ser Val Gln Ser Ser Ile Gly Asn Leu Ile Val Ala Ile Lys
            20
                                25
Ser Val Gln
        35
<210> 1478
<211> 35
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1478
Lys Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val
                                   10
Gln Ser Ser Ile Gly Asn Leu Ile Val Ala Ile Lys Ser Val Gln Asp
                                25
            20
Tyr Val Asn
       35
<210> 1479
<211> 35
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1479 Leu Lys Glu Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln 5 Ser Ser Ile Gly Asn Leu Ile Val Ala Ile Lys Ser Val Gln Asp Tyr 25 20 Val Asn Lys 35 <210> 1480 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1480 Ala Ile Arg Asp Thr Asn Lys Ala Val Gln Ser Val Gln Ser Ser Ile 5 Gly Asn Leu Ile Val Ala Ile Lys Ser Val Gln Asp Tyr Val Asn Lys 25 20 Glu Ile Val 35 <210> 1481 <211> 47 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1481 Thr Trp Gln Glu Trp Glu Arg Lys Val Asp Phe Leu Glu Glu Asn Ile 5 Thr Ala Leu Leu Glu Glu Ala Gln Ile Gln Gln Glu Lys Asn Met Tyr 20 25

Glu Leu Gln Lys Leu Asn Ser Trp Asp Val Phe Gly Asn Trp Phe

35 40 45

<210> 1482

<211> 35

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1482

Trp Gln Glu Trp Glu Arg Lys Val Asp Phe Leu Glu Glu Asn Ile Thr
1 5 10 15

Ala Leu Leu Glu Glu Ala Gln Ile Gln Gln Glu Lys Asn Met Tyr Glu 20 25 30

Leu Gln Lys

35

<210> 1483

<211> 35

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 Peptide

<400> 1483

Gln Glu Trp Glu Arg Lys Val Asp Phe Leu Glu Glu Asn Ile Thr Ala 1 5 10 15

Leu Leu Glu Glu Ala Gln Ile Gln Gln Glu Lys Asn Met Tyr Glu Leu 20 25 30

Gln Lys Leu

35

<210> 1484

<211> 35

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: Synthetic

<220>

Peptide

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<400> 1484
 Glu Trp Glu Arg Lys Val Asp Phe Leu Glu Glu Asn Ile Thr Ala Leu
 Leu Glu Glu Ala Gln Ile Gln Glu Lys Asn Met Tyr Glu Leu Gln
                                  25
 Lys Leu Asn
          35
 <210> 1485
 <211> 35
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
       Peptide
  <400> 1485
 Trp Glu Arg Lys Val Asp Phe Leu Glu Glu Asn Ile Thr Ala Leu Leu
 Glu Glu Ala Gln Ile Gln Gln Glu Lys Asn Met Tyr Glu Leu Gln Lys
                                  25
  Leu Asn Ser
          35
<210> 1486
  <211> 35
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Synthetic
       Peptide
  <400> 1486
  Glu Arg Lys Val Asp Phe Leu Glu Glu Asn Ile Thr Ala Leu Leu Glu
                    5
                                     10
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Glu Ala Gln Ile Gln Gln Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu 25 20 Asn Ser Trp 35 <210> 1487 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1487 Arg Lys Val Asp Phe Leu Glu Glu Asn Ile Thr Ala Leu Leu Glu Glu 10 Ala Gln Ile Gln Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu Asn 25 Ser Trp Asp 35 <210> 1488 <211> 35 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1488 Lys Val Asp Phe Leu Glu Glu Asn Ile Thr Ala Leu Leu Glu Glu Ala · 10 Gln Ile Gln Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu Asn Ser 25 Trp Asp Val 35

502

<210> 1489

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<211> 35
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 Ile Gln Gln Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu Asn Ser Trp
                                  25
 Asp Val Phe
          35
 <210> 1490
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                                      10
 Gln Gln Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu Asn Ser Trp Asp
                                  25
· Val Phe Gly
         35
 <210> 1491
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  <223> Description of Artificial Sequence: Synthetic
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<400> 1491 Phe Leu Glu Glu Asn Ile Thr Ala Leu Leu Glu Glu Ala Gln Ile Gln 10 Gln Glu Lys Asn Met Tyr Glu Leu Gln Lys Leu Asn Ser Trp Asp Val 20 25 Phe Gly Asn 35 <210> 1492 <211> 35 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1492 Pro Asp Ala Val Tyr Leu His Arg Ile Asp Leu Gly Pro Pro Ile Ser 5 Leu Glu Arg Leu Asp Val Gly Thr Asn Leu Gly Asn Ala Ile Ala Lys 20 25 Leu Glu Asp 35 <210> 1493 <211> 34 <212> PRT <213> Altificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1493 Leu Glu Arg Leu Asp Val Gly Thr Asn Leu Gly Asn Ala Ile Ala Lys 5 10 15 Leu Glu Ala Lys Glu Leu Leu Glu Ser Ser Asp Gln Ile Leu Arg Ser - 20 25

504

Met Lys

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<210> 1494
<211> 34
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Leu His Arg Ile Asp Leu Gly Pro Pro Ile Ser Leu Glu Arg Leu Asp
                  5
Val Gly Thr Asn Leu Gly Asn Ala Ile Ala Lys Leu Glu Ala Lys Glu
                                 25
Leu Leu
<210> 1495
<211> 34
<212> PRT
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His Arg Ile Asp Leu Gly Pro Pro Ile Ser Leu Glu Arg Leu Asp Val
                                     10
                 5
Gly Thr Asn Leu Gly Asn Ala Ile Ala Lys Leu Glu Ala Lys Glu Leu
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Leu Glu
 <210> 1496
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<220>

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    Arg Ile Asp Leu Gly Pro Pro Ile Ser Leu Glu Arg Leu Asp Val Gly
      1 5
    Thr Asn Leu Gly Asn Ala Ile Ala Lys Leu Glu Ala Lys Glu Leu Leu
                                  25
                20
    Glu Ser
    <210> 1497
    <211> 34
    <212> PRT
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    <400> 1497
    Ile Asp Leu Gly Pro Pro Ile Ser Leu Glu Arg Leu Asp Val Gly Thr
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    Asn Leu Gly Asn Ala Ile Ala Lys Leu Glu Ala Lys Glu Leu Leu Glu
    Ser Ser
<210> 1498
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    <212> PRT
    <213> Artificial Sequence
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    <400> 1498
    Asp Leu Gly Pro Pro Ile Ser Leu Glu Arg Leu Asp Val Gly Thr Asn
                                      10
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Leu Gly Asn Ala Ile Ala Lys Leu Glu Ala Lys Glu Leu Leu Glu Ser 20 25 30

Ser Asp

<210> 1499

<211> 34

<212> PRT

<213> Artificial Sequence

<220×

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1499

Leu Gly Pro Pro Ile Ser Leu Glu Arg Leu Asp Val Gly Thr Asn Leu

1 5 10 15

Gly Asn Ala Ile Ala Lys Leu Glu Ala Lys Glu Leu Leu Glu Ser Ser 20 25 30

Asp Gln

<210> 1500

<211> 34

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1500

Gly Pro Pro Ile Ser Leu Glu Arg Leu Asp Val Gly Thr Asn Leu Gly
1 5 10 15

Asn Ala Ile Ala Lys Leu Glu Ala Lys Glu Leu Leu Glu Ser Ser Asp 20 25 30

Gln Ile

<210> 1501

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<211> 34
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                  5
                                     10
Ala Ile Ala Lys Leu Glu Ala Lys Glu Leu Leu Glu Ser Ser Asp Gln
                                 25
Ile Leu
<210> 1502
<211> 34
<212> PRT
<213> Artificial Sequence
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<400> 1502
Pro Ile Ser Leu Glu Arg Leu Asp Val Gly Thr Asn Leu Gly Asn Ala
                                     10
Ile Ala Lys Leu Glu Ala Lys Glu Leu Leu Glu Ser Ser Asp Gln Ile
                                 25
Leu Arg
<210> 1503
<211> 34
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<400> 1503 Ser Leu Glu Arg Leu Asp Val Gly Thr Asn Leu Gly Asn Ala Ile Ala 5 10 Lys Leu Glu Ala Lys Glu Leu Leu Glu Ser Ser Asp Gln Ile Leu Arg 20 25 Ser Met <210> 1504 <211> 34 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1504 Leu Glu Arg Leu Asp Val Gly Thr Asn Leu Gly Asn Ala Ile Ala Lys 5 10 Leu Glu Ala Lys Glu Leu Leu Glu Ser Ser Asp Gln Ile Leu Arg Ser 20 25 30 Met Lys <210> 1505 <211> 22 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1505 Tyr Leu Cys Glu Phe Cys Leu Lys Tyr Gly Arg Ser Leu Lys Cys Leu 10 Gln Arg His Leu Thr Lys

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<210> 1506
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Ala Ser Thr Thr Thr Asn Tyr Thr
1 5
<210> 1507
<211> 8
<212> PRT
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<400> 1507
Ala Ser Thr Thr Thr Asn Tyr Thr
1 5
<210> 1508
<211> 8
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
    Peptide
<400> 1508
Ser Gln Asn Tyr Pro Ile Val Gln
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<210> 1509
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1509
Glu Leu Asp Lys Trp Ala
                5
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<210> 1510
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<400> 1510
Arg Gly Val Val Asn Ala Ser Ser Arg Leu Ala
                                    10
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                 5
<210> 1511
<211> 10
<212> PRT
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<400> 1511
Arg Gly Arg Arg Gln Pro Ile Pro Lys Ala
<210> 1512
<211> 23
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1512
Leu Thr Thr Gly Ser Val Val Ile Val Gly Arg Ile Ile Leu Ser Gly
                  5
                                    10
                                                         15
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Arg Pro Ala Val Val Pro Asp 20

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<210> 1513
<211> 14
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1513
Gly Ser Val Val Ile Val Gly Arg Ile Ile Leu Ser Gly Arg
       5
<210> 1514
<211> 27
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Pro Leu Gly Phe Phe Pro Asp His Gln Leu Asp Pro Ala Phe Gly Ala
Asn Ser Asn Asn Pro Asp Trp Asp Phe Asn Pro
            20
<210> 1515
<211> 26
<212> PRT
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<400> 1515
Met Gln Trp Asn Ser Thr Thr Phe His Gln Thr Leu Gln Asp Pro Arg
                                    10
                                                       15
                  5
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Val Arg Gly Leu Tyr Phe Pro Ala Gly Gly
            20
                              25
<210> 1516
<211> 11
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1516
Met Gln Trp Asn Ser Thr Ala Phe His Gln Thr
     5
<210> 1517
<211> 8
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1517
Tyr Gly Ala Val Val Asn Asp Leu
                5
1
<210> 1518
<211> 21
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1518
Cys Thr His Gly Ile Arg Pro Val Val Ser Thr Gln Leu Leu Leu Asn
                5
                                  10
Gly Ser Leu Ala Glu
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<210> 1519
<211> 7
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1519
Ser Glu Asn Tyr Pro Ile Val
<210> 1520
<211> 7
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1520
Lys Ala Arg Val Phe Glu Ala
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                5
<210> 1521
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1521
Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe Val Thr Ile Gly Lys
                 5
                            10
<210>_ 1522
<211> 8
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1522
Ala Ser Thr Thr Thr Asn Tyr Thr
1 5
<210> 1523
<211> 8
<212> PRT
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     Peptide
<400> 1523
Ala Ser Thr Thr Thr Asn Tyr Thr
1 5
<210> 1524
<211> 15
<212> PRT
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     Peptide
<400> 1524
Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe Val Thr Ile Gly Lys
                          10
<210> 1525
<211> 38
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1525
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Asn Asn Leu Leu Arg Ala Ile Glu Ala Gln Gln His Leu Leu Gln Leu
 1 5
                                  10
Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile Leu Ala Val Glu
            20
                               25
Arg Tyr Leu Lys Asp Gln
        35
<210> 1526
<211> 10
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1526
Arg Gly Pro Gly Arg Ala Phe Val Thr Ile
1 5
<210> 1527
<211> 3
<212> PRT
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     Peptide
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Phe Phe Gly
1
<210> 1528
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
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<400> 1528

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Tyr Thr Ser Leu Ile His Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln
 1 5
                                 10
Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
Trp Asn Trp Phe
        35
<210> 1529
<211> 2
<212> PRT
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<400> 1529
Thr Gln
 1
<210> 1530
<211> 6
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1530
Trp His Trp Leu Gln Leu
1 5
<210> 1531
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1531
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Gly Pro Gly Ala Gly
1 5
<210> 1532
<211> 2
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1532
His Gly
1
<210> 1533
<211> 5
<212> PRT
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     Peptide
<400> 1533
Phe Val Phe Leu Met
<210> 1534
<211> 3
<212> PRT
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     Peptide
<400> 1534
Thr Ser Lys
1
<210> 1535
<211> 3
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<212> PRT
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  <400> 1535
  Lys His Gly
   1
  <210> 1536
  <211> 16
  <212> PRT
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        Peptide
  <400> 1536
  Lys Asn Arg Trp Glu Asp Pro Gly Lys Gln Leu Tyr Asn Val Glu Ala
                                      10
  <210> 1537
  <211> 10
  <212> PRT
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  <223> Description of Artificial Sequence: Synthetic
        Peptide
  <400> 1537
  Leu Arg Ala His Ala Val Asp Val Asn Gly
                    5
  <210> 1538
. <211> 14
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: Synthetic
        Peptide
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<400> 1538
 Trp Ser Lys Met Asp Gln Leu Ala Lys Glu Leu Thr Ala Glu
 1 5
                                 10
 <210> 1539
 <211> 4
 <212> PRT .
 <213> Artificial Sequence
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      Peptide
 <400> 1539
 Thr Pro Arg Lys
  1
 <210> 1540
 <211> 12
 <212> PRT
 <213> Artificial Sequence
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 <223> Description of Artificial Sequence: Synthetic
      Peptide
 <400> 1540
 Gly Glu Leu Gln Asn Gln Leu Ile Arg Lys Ser Asn
  1 5 10
 <210> 1541
 <211> 17
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      Peptide
 <400> 1541
 Gly Glu Tyr Gln Lys Met Leu Asn Leu Arg Ala Glu Val Lys Lys Asn
 Ala
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<210> 1542
<211> 9
<212> PRT
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     Peptide
<400> 1542
Pro Phe Cys Asn Ala Phe Thr Gly Cys
                5
<210> 1543
<211> 8
<212> PRT
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     Peptide
<400> 1543
Glu Lys Ala His Asp Gly Gly Arg
                5
<210> 1544
<211> 14
<212> PRT
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      Peptide
<400> 1544
Pro Phe Thr Arg Asn Tyr Tyr Val Arg Ala Val Leu His Leu
<210> 1545
<211> 39
<212> PRT
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       Peptide
 <400> 1545
 Ala Pro Arg Leu Pro Gln Cys Gln Gly Asp Asp Gln Glu Lys Cys Leu
 Cys Asn Lys Asp Glu Cys Pro Pro Gly Gln Cys Arg Phe Pro Arg Gly
                        25
 Asp Ala Asp Pro Tyr Cys Glu
         35
 <210> 1546
 <211> 8
 <212> PRT
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       Peptide
 <400> 1546
 Lys Gly Asp Glu Glu Ser Leu Ala
 <210> 1547
 <211> 9
 <212> PRT
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       Peptide
 <400> 1547
 Trp Ala Gly Gly Asp Ala Ser Gly Glu
  1
                   5
 <210> 1548
 <211> 20
 <212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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Gln Ala Thr Val Gly Asp Val Asn Thr Asp Arg Pro Gly Leu Leu Asp
                                    10
Leu Lys Tyr Tyr
            20
<210> 1549
<211> 17
<212> PRT
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      Peptide
<400> 1549
Thr Lys Arg Arg Ala Ile Gly Phe Lys Leu Ala Glu Ala Val Lys
                  5
                                    10
                                                         15
Cys
<210> 1550
<211> 8
<212> PRT
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      Peptide
<400> 1550
Ser Ile Ile Asn Phe Glu Lys Leu
 1
                 5
<210> 1551
<211> 6
<212> PRT
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     Peptide
<400> 1551
Phe Gln Val Val Cys Gly
 1
      5
<210> 1552
<211> 6
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1552
Ala Arg Met Ala Pro Glu
                 5
<210> 1553
<211> 4
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1553
Gly Gln Pro Arg
 1
<210> 1554
<211> 3
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1554
Thr Val Leu
<210> 1555
<211> 49
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1555
Leu Leu Ser Lys Arg Gly His Cys Pro Arg Ile Leu Phe Arg Cys Pro
                 5
Leu Ser Asn Pro Ser Asn Lys Cys Trp Arg Asp Tyr Asp Cys Pro Gly
            20
                               25
Val Lys Lys Cys Cys Glu Gly Phe Cys Gly Lys Asp Cys Leu Tyr Pro
                           40
Lys
<210> 1556
<211> 10
<212> PRT
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     Peptide
<400> 1556
Gly Phe Asp Leu Asn Gly Gly Gly Val Gly
                5
                                   10
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<210> 1557
<211> 18
<212> PRT
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<220>
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Ala Val Gln Ser Lys Pro Pro Ser Lys Arg Asp Pro Pro Lys Met Gln
                5
                                  10
Thr Asp
<210> 1558
<211> 4
<212> PRT
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     Peptide
<400> 1558
Thr Lys Pro Arg
 1
<210> 1559
<211> 11
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1559
Tyr Leu Asn Phe Thr Pro Asn Trp Gly Thr Tyr
 1 5 10
<210> 1560
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     Peptide
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<400> 1560
Asp Leu Trp Gln Lys
<210> 1561
<211> 17
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     Peptide
<400> 1561
Trp Tyr Glu Pro Ile Tyr Leu Gly Gly Val Phe Gln Leu Glu Lys Gly
                      10
Asp
<210> 1562
<211> 39
<212> PRT
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     Peptide
<400> 1562
Ala Pro Arg Leu Pro Gln Cys Gln Gly Asp Gln Glu Lys Cys Leu Cys
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Asn Lys Asp Glu Cys Pro Pro Gly Gln Cys Arg Phe Pro Arg Gly Asp
        20 25
Ala Asp Pro Tyr Cys Glu Asp
        35
<210> 1563
<211> 4
<212> PRT
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<220>
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<223> Description of Artificial Sequence: Synthetic
     Peptide
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Thr Lys Pro Arg
 1
<210> 1564
<211> 6
<212> PRT
<213> Artificial Sequence
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     Peptide
<400> 1564
Glu Glu Val Val Ala Cys
 1
<210> 1565
<211> 4
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1565
Ser Asp Lys Pro
 1
<210> 1566
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1566
Arg Phe Trp Ile Asn Lys
 1
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<210> 1567
<211> 13
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1567
Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly
      5
                                  10
<210> 1568
<211> 1
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1568
Leu
1
<210> 1569
<211> 5
<212> PRT
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     Peptide
<400> 1569
Asp Asp Asp Asp
 1
<210> 1570
<211> 6
<212> PRT
<213> Artificial Sequence
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<220>
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 1570
 Asp Asp Asp Asp Asp
  1
                   5
 <210> 1571
 <211> 8
 <212> PRT
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       Peptide
 <400> 1571
 Asn Pro Asn Ala Asn Pro Asn Ala
 <210> 1572
 <211> 8
 <212> PRT
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 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 1572
 Val Ala Ile Thr Val Leu Val Lys
                  5
 <210> 1573
 <211> 6
 <212> PRT
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  <223> Description of Artificial Sequence: Synthetic
       Peptide
  <400> 1573
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Val Gly Val Arg Val Arg
   1
  <210> 1574
  <211> 4
  <212> PRT
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        Peptide
  <400> 1574
  Val Ile His Ser
   1
  <210> 1575
  <211> 5
  <212> PRT
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  <223> Description of Artificial Sequence: Synthetic
        Peptide
  <400> 1575
  Val Pro Asp Pro Arg
  <210> 1576
  <211> 4
  <212> PRT
  <213> Artificial Sequence
                           <220>
  <223> Description of Artificial Sequence: Synthetic
        Peptide
  <400> 1576
  Val Thr Cys Gly
   1
<210> 1577
  <211> 3
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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 1577
Arg Ser Arg
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<210> 1579
<211> 6
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Asp His Ala Arg Trp Lys
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<210> 1580
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 1580

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Pro Gln Asp Pro Gln Asp Leu
       5
<210> 1581
<211> 6
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Gln His Phe Arg Trp Gly
<210> 1582
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Asn Gln Glu Gln Val Ser Pro Leu Thr Leu Leu Lys Leu Gly Asn Gln
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Glu Pro Gly
<210> 1583
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Leu Ser Ala Leu Ser Leu Asp Glu Pro Phe Ile Gln Lys Asp Val Glu
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5 10 15 1 Leu Arg Ile Met <210> 1584 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1584 Ala Pro Glu Ala Gln Val Ser Val Gln Pro Asn Phe Gln Gln Asp 5 10 <210> 1585 <211> 27 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Peptide <400> 1585 Glu Tyr Gly Gly Thr Lys Val Leu Asp Asp Lys Asp Tyr Phe Leu Phe 15 5 10 Arg Asp Gly Asp Ile Leu Gly Lys Tyr Val Asp <210> 1586 <211> 16 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic Peptide <400> 1586 Lys Ala Tyr Ile Asn Lys Val Glu Glu Leu Lys Lys Lys Tyr Gly Ile

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1 5 10
<210> 1591
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<400> 1591
Ala Leu Thr Asp Phe Phe Arg
1 5
<210> 1592
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<223> Description of Artificial Sequence: Synthetic
     Peptide
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Gln Ala Ile Gly Leu Met Gly Tyr Arg Leu Ser Pro Gln Thr Leu Thr
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Thr Ile Val Lys
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<210> 1593
<211> 17
<212> PRT
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Met Gly Phe Asn Ala Phe Lys Glu Leu Trp Ala Ala Leu Asn Ala Trp
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Lys
<210> 1594
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      Peptide
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Glu Val Gln Leu Val Glu Ser Gly Val Gly Leu Val Gln Pro Gly Asp
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                                                         15
<210> 1595
<211> 17
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<400> 1595
Glu Leu Asp Ala Lys Ile Pro Ser Thr Gly Asp Ala Thr Glu Trp Arg
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Asn
<210> 1596
<211> 20
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Val Thr Ser Ala
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<210> 1597
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      Peptide
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Gly Lys Glu Ile Leu Val Gly Asp Val Gly Gln Thr Val Asp Asp Pro
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Tyr Ala Thr Phe
<210> 1598
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Glu Leu Ser Leu Ala Gly Asn Glu Leu Gly Asp Glu Gly Ala Arg
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<210> 1599
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Met Phe Ile Val Asn Thr Asn Val Pro Arg Ala Ser Val Pro Asp Gly
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Phe Leu Ser Glu Leu Thr
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<210> 1600
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Phe Gly Ser Phe Phe Thr Leu Asn Leu Phe Ile Gly Ile Ile Ile Asp
Asn Phe Trp Gln Gln Lys Lys Lys Leu Gly Gly Lys Asp Ile Phe Met
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Thr Glu Glu
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<210> 1601
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      Peptide
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Gly Leu Pro Gly Arg Asp Gly Arg Asp Gly Arg Glu Gly Phe Arg Gly
                                                          15
Glu Glu Gly Asp Pro Gly Leu Pro Gly Ala Ala
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<210> 1602
<211> 21
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     Peptide
<400> 1602
Cys Lys Thr Cys Gln Arg Lys Phe Ser Arg Ser Gly His Leu Lys Thr
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His Thr Arg Thr His
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<210> 1603
<211> 15
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     Peptide
<400> 1603
His Val Ser Ser Glu Ala Phe Arg Met Cys Asp Val Cys Leu Glu
     5
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                                         15
<210> 1604
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Pro His Ser Arg Asn
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<210> 1605
<211> 5
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<223> Description of Artificial Sequence: Synthetic
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Pro His Ser Cys Asn
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<210> 1606
<211> 28
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      Peptide
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Met Gly Pro Gly Ala Pro Phe Ala Arg Val Gly Trp Pro Leu Pro Leu
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Leu Val Val Met Ala Gly Val Ala Pro Val Trp Ala
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<210> 1607
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Met Val Lys Ser His Ile Gly Ser Trp Ile Leu Val Leu Phe Val
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                                                          15
<210> 1608
<211> 9
<212> PRT
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 Cys Ser Leu Pro Gly Ser Ala Ala Ala
  1 5
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 <211> 15
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 Arg Gly Leu Lys Arg Gln Ser Asp Glu Arg Lys Arg Asp Arg Glu
 <210> 1610
 <211> 57
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       Peptide
 Gly Arg Ser His Met Val Leu Asn Ser Ala Leu Glu Gly Ala Arg Gly
1 5 10
                                       1.5
 Gly Pro Gly Gly Glu Glu Ile Pro Glu Arg Phe Ser Ile Pro Glu Leu
                               25
             20
 Gln Trp Met Leu Ser Asn Ala Glu Leu Ala Pro Val Gln Ala Asp Glu
         35
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 Pro Pro Gln Ser Arg Met Asp Leu Val
      50
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Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala Asp Arg Ala
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Ala Val Met
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<210> 1612
<211> 15
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Met Val Lys Ser His Ile Gly Ser Trp Ile Leu Val Leu Phe Val
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1 5
                       10
<210> 1613
<211> 29
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<400> 1613
Leu Arg Asp Leu Val Ser Tyr Cys Arg Ala Arg Gly Lys Gly Arg Glu
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Arg Met Asn Gly Thr Arg Lys Gly His Leu Leu Tyr Met
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<210> 1614

<211> 33

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Peptide

<400> 1614

Gly Lys Arg Ser Ser Pro Glu Thr Leu Ile Ser Asp Leu Leu Met Arg

1 5 10 15

Glu Ser Thr Glu Asn Val Pro Arg Thr Arg Leu Glu Asp Pro Ala Met 20 25 30

Trp

<210> 1615

<211> 30

<212> PRT

<213> Artificial Sequence

<220>

<400> 1615

Asp Val Glu Pro Leu Leu Gly Phe Leu Ser Pro Lys Ser Gly Gln Glu
1 5 10 15

Asn Glu Val Asp Asp Phe Pro Tyr Lys Gly Gln Gly Glu Leu 20 25 30

<210> 1616

<211> 20

<212> PRT

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